



Ingenieur fakultät Bau Geo Umwelt  
Signalverarbeitung in der Erdbeobachtung

# **Advances in Meter-resolution Multipass Synthetic Aperture Radar Interferometry**

Yuanyuan Wang, M.Sc.

Vollständiger Abdruck  
der von der Ingenieur fakultät Bau Geo Umwelt  
der Technischen Universität München  
zur Erlangung des akademischen Grades eines  
Doktor-Ingenieurs (Dr.-Ing.)  
genehmigten Dissertation

Vorsitzender: Univ.-Prof. Dr.-Ing. habil. Florian Seitz

Prüfer der Dissertation:

1. Univ.-Prof. Dr.-Ing. habil. Xiaoxiang Zhu
2. Univ.-Prof. Dr.-Ing. habil. Richard Bamler
3. Univ.-Prof. Dr.-Ing. Uwe Sörgel  
Technische Universität Darmstadt

Die Dissertation wurde am 19.10.2015 bei der Technischen Universität München eingereicht und durch die Ingenieur fakultät Bau Geo Umwelt am 04.11.2015 angenommen.





# Abstract

**S**ynthetic aperture radar (SAR) interferometry is the technique to use SAR as interferometer, to measure the phase difference caused by topography or object displacement. Multipass SAR interferometry (InSAR) is so far the only method for assessing long-term millimetre-level deformation over large areas from space on an imaging basis. Very high resolution multipass InSAR techniques have made substantial development in the last decade in monitoring individual building using persistent scatterer interferometry (PSI) and SAR tomography (TomoSAR), nonurban area using small baseline subset (SBAS) and SqueeSAR, and so on.

To prepare for future SAR missions which will have higher resolution, larger coverage, and greater data volume, this thesis addresses the following four aspects of multipass InSAR techniques: computational efficiency, information fusion, contextual awareness, and statistical robustness.

TomoSAR is the most competent InSAR method for urban area monitoring. But it is much more computationally intensive than any other multipass InSAR methods. To this end, an efficient multi-dimensional TomoSAR algorithm for urban area was developed, by *integrating tomographic SAR inversion and the well-known PSI*. The results of PSI were used for a pre-classification of single and double scatterers, and also used as prior in the TomoSAR reconstruction. Real data experiments show the proposed approach obtains results comparable to the one obtained by the most computationally expensive tomographic SAR algorithms (e.g. SL1MMER), and saves computational time by a factor of fifty.

*Multi-aspect InSAR point clouds fusion* is required for a complete monitoring of entire city, due to the SAR side-looking geometry. In this thesis, a robust algorithm, namely "L-shape detection & matching", is proposed, especially for fusing two point clouds from cross-heading orbits, i.e. ascending and descending. The main idea of this algorithm is finding and matching the theoretically exact point correspondence which is the end positions of façades where the two point clouds close. Practical experiment shows the proposed method achieves sub-meter consistency with the state-of-the-art, and is much more computationally efficient.

The development of *a semantic urban infrastructure monitoring algorithm by fusing InSAR and optical images* was followed after the point cloud fusion. The attributes derived from optical images, e.g. colour, classification label, are transferred to the InSAR point cloud for a semantic level analysis of the deformation signal. The key lies on a strict 3-D geometric co-registration of SAR and optical images by reconstructing and matching the 3-D point clouds derived from the two types of images. Examples on bridges and railway monitoring are demonstrated.

*Robust InSAR optimization* is crucial for future SAR data, because of the unprecedented high resolution brings different observation statistics and much more dynamic interferometric phase. The proposed robust InSAR optimization (RIO) framework answers two open questions in multipass InSAR: (1) How to optimally treat images with a large phase error, e.g., due to unmodeled motion phase, uncompensated atmospheric phase, etc.? And (2) How to estimate the covariance matrix of a non-Gaussian complex InSAR multivariate, particularly those with nonstationary phase signals? For the former question, RIO employs a robust M-estimator to effectively down-weight these images, and for the latter question, a new method — the rank M-Estimator — is proposed. Simulated and real data experiments demonstrated that substantial improvement can be achieved in terms of the variance of estimates, comparing to the state-of-the-art estimators for persistent and distributed scatterers. The proposed framework can be easily extended to other multipass InSAR techniques.

The abovementioned algorithms were tested using TerraSAR-X data of various test sites,

especially for the efficient TomoSAR algorithm and the RIO framework.

**Keywords**

Synthetic aperture radar interferometry, robust estimation, covariance matrix, distributed scatterer, persistent scatterer, SAR tomography, deformation monitoring, semantic interpretation, InSAR point cloud, data fusion

# Zusammenfassung

**S**ynthetisch Apertur Radar-Interferometrie ist eine Technologie um die Phase die durch die Topographie oder die Verschiebung von Objekten entsteht zu messen. Multipass-SAR-Interferometrie ist bis heute das einzige bildgebende Verfahren um über weite Gebiete langfristig millimetergenau Deformation mit Satelliten zu messen. Multipass SAR-Techniken für sehr hohe Auflösungen haben in der letzten Dekade eine signifikante Entwicklung erfahren, sei es bei der Beobachtung von einzelnen Gebäuden mit Persistenter Streuer Interferometrie (PSI) und SAR-Tomographie (TomoSAR), bei nichturbanen Gebieten mittels Small Baseline Subset (SBAS) und SqueeSAR, und so weiter.

Um für zukünftige SAR Missionen bereit zu sein, welche eine höhere Auflösung, größere Abdeckung und ein größeres Datenvolumen besitzen, behandelt diese Dissertation die folgenden vier Aspekte von Multipass-InSAR-Techniken: Recheneffizienz, Fusion von Information, Berücksichtigung von Zusammenhängen und robuste Schätzverfahren.

TomoSAR ist die leistungsfähigste InSAR-Methode für die Beobachtung von urbanen Gebieten. Allerdings ist es weitaus rechenintensiver als alle anderen Multipass-InSAR-Methoden. Daher wurde ein effizienter Ansatz für mehrdimensionale tomographische SAR-Inversion für urbane Gebiete entwickelt, indem *die tomographischer SAR-Inversion in die bekannte PSI integriert* wurde. Die Ergebnisse der PSI wurden für eine Vorklassifizierung in Einzel- und Doppelstreuer benutzt und fungierten als *a priori*-Wissen für die TomoSAR-Rekonstruktion. Versuchsläufe mit Realdaten zeigen, dass der vorgeschlagene Ansatz Ergebnisse liefert die vergleichbar sind mit denen der rechenintensivsten tomographischen SAR-Algorithmen (z.B. SLIMMER) und die Rechenzeit um den Faktor fünfzig verkürzt.

*Die Fusion von Multi-aspect InSAR-Punktwolken* ist aufgrund der seitlich schauenden Geometrie notwendig um eine ganze Stadt vollständig zu beobachten. In dieser Dissertation wird ein robuster Algorithmus — L-Form Detektion und Zuordnung — zur Fusion von Punktwolken, vor allem von cross-heading Orbits, das heißt ascending und descending, vorgeschlagen. Die Grundidee dieses Algorithmus ist das Finden und die Zuordnung der theoretisch exakten Punktkorrespondenzen, welche die Endpunkte von Fassaden sind an denen zwei Punktwolken zusammenlaufen. Praktische Experimente zeigen, dass die vorgeschlagene Methode Übereinstimmungen von unter einem Meter mit Methoden auf dem Stand der Technik erreicht, bei gleichzeitig weitaus effizienterer Nutzung der Rechenzeit.

Die Entwicklung eines Algorithmus für die *semantische Beobachtung von urbaner Infrastruktur mittels Fusion von InSAR und optischen Bildern* folgte nach der Punktwolken-Fusion. Anschließend werden die vom optischen Bild erhaltenen Attribute — Farbe, Klassifizierung, usw. — in die InSAR Punktwolke für eine Semantikanalyse des Deformationssignals übertragen. Entscheidend ist die strenge 3-D geometrische Koregistrierung von SAR-Bildern und optischen Bildern durch die Rekonstruktion und die Zuordnung der 3-D Punktwolken welche von den zwei Typen von Bildern erzeugt wurden. Der vorgeschlagene Ansatz wird anhand von Beispieldaten mit Brücken und Eisenbahnschienen gezeigt.

*Robuste InSAR Optimierung* ist entscheidend für zukünftige SAR-Daten, da die bisher noch nie dagewesene hohe Auflösung neue Beobachtungsstatistiken und eine weitaus dynamischere Phase mit sich bringt. Das vorgeschlagene robuste InSAR-Optimierungsframework (RIO) beantwortet zwei offene Fragen für Multipass-InSAR: (1) Wie sollen Bilder mit großen Phasenfehlern optimal behandelt werden, z.B. aufgrund nicht im Modell berücksichtigter durch Bewegung verursachte Phasenänderung, nicht kompensierter atmosphärischer Phase, usw.? (2) Wie kann die Kovarianzmatrix für nicht-Gaussche, komplexe, mehrdi-

mensionale InSAR Zufallsvariable geschätzt werden, insbesondere die mit nicht stationären Phasensignalen? Um die erste Frage zu adressieren setzt RIO einen robusten M-Schätzer ein, welcher Bildern mit großen Phasenfehlern ein geringeres Gewicht gibt, für die zweite Frage wird eine neue Methode — der Rang M-Schätzer — vorgeschlagen, der kein phase flattening benötigt. Versuche zeigen, dass erhebliche Verbesserungen bezüglich der Varianz der Schätzung erreicht werden können, im Vergleich zu Schätzern auf dem Stand der Technik für persistente und verteilte Streuer. Das vorgeschlagene Framework kann leicht für andere InSAR Techniken erweitert werden.

Die oben erwähnten Algorithmen wurden mit TerraSAR-X für mehrere Testgebiete getestet, insbesondere gilt dies für den effizienten TomoSAR-Algorithmus und das RIO Framework.

# List of Abbreviations

Abbreviation	Description
2-D	two-dimensional
3-D	three-dimensional
AD	Anderson-Darling
ADI	amplitude dispersion index
AIC	Akaike information criterion
APS	atmospheric phase screen
BIC	Bayesian information criterion
BoW	bag of words
CCG	complex circular Gaussian
CDF	cumulative distribution function
CRLB	Cramér-Rao lower bound
CS	compressive sensing
D-InSAR	differential SAR interferometry
DoF	degree of freedom
DS	distributed scatterer
D-TomoSAR	differential SAR tomography
DEM	digital elevation model
GPSR	gradient projection for sparse reconstruction
HPR	hidden point removal
ICP	iterative closest point
i.i.d.	independent and identical distribution
InSAR	SAR interferometry
KS	Kolmogorov-Smirnov
LAMBDA	least square ambiguity decorrelation

Abbreviation	Description
LOS	line-of-sight
MAP	maximum a posteriori
MLE	maximum likelihood estimator
MUSIC	multiple signal classification
NLS	nonlinear least squares
PDF	probability density function
PCA	principle component analysis
PS	persistent scatterer
PSI	persistent scatterer interferometry
RANSAC	random sample consensus
RIO	robust InSAR optimization
RME	rank M-estimator
RSS	residual sum of squares
SAR	synthetic aperture radar
SBAS	small baseline subset
SCM	sign covariance matrix
SfM	structure from motion
SL1MMER	scale-down by $L_1$ norm minimization, model selection, and estimation reconstruction
SLC	single look complex
SNR	signal to noise ratio
SVD	singular value decomposition
SVM	support vector machine
TerraSAR-X	TS-X
TomoSAR	SAR tomography
UTM	universal transverse Mercator

# List of Symbols

Symbol	Description
$g$	single SAR pixel value
$I$	intensity
$A$	amplitude
$\phi$	unambiguous (absolute) phase
$\varphi$	wrapped phase
$R$	range distance
$R_s$	nominal range distance
$\lambda$	wavelength
$\theta$	incidence angle
$B_{\perp}$	perpendicular baseline
$h$	height
$v$	linear deformation rate
$s$	elevation
$\Delta s$	elevation extent
$\gamma$	coherence
$\gamma(s)$	reflectivity profile along elevation direction
$\Upsilon$	discrete reflectivity profile
$L$	number of elements in $\gamma$
$L_s$	number of elements in $\gamma$ required for sparse reconstruction
$K$	number of scatterers
$K_0(\cdot)$	modified Bessel function of the second kind
$\text{Li}_2$	dilogarithm
$N$	number of images
$M$	number of samples
$\mathbf{g}$	multipass / multibaseline SAR pixel values
$g_n$	$n$ th element of $\mathbf{g}$
$\mathbf{g}_m$	$m$ th neighbouring sample of $\mathbf{g}$
$\hat{\mathbf{r}}$	estimated complex rank of $\mathbf{g}$

Symbol	Description
$\theta$	phase history parameters
$\mathbf{a}$	integer $2\pi$ phase ambiguities
$\mathbf{C}$	complex covariance matrix
$\hat{\mathbf{C}}_k$	covariance estimate in the $k$ th iteration
$\mathbf{C}_{\gamma\gamma}$	prior covariance matrix of $\gamma$
$\mathbf{R}$	SAR discrete imaging model matrix
$\Phi(\theta)$	diagonal matrix of the modelled phase
$\mathbf{W}$	diagonal weighting matrix
$\sigma$	standard deviation
$F$	distribution function
$x$	generic variable
$\varepsilon$	residual vector
$\varepsilon_i$	$i$ th element of $\varepsilon$
$\alpha$	regularization parameter
$\nu$	degree of freedom
$\eta_{RE}$	relative efficiency
$H_0$	null hypothesis (no scatterer)
$H_{1,2}$	alternative hypothesis (single or double scatterer)
$P_0$	prior probability of the null hypothesis
$P_{1,2}$	prior probability of the alternative hypothesis
$\rho(\cdot)$	loss function
$\psi(\cdot)$	influence function
$w(\cdot)$	weighting function
$p(\cdot \cdot)$	likelihood function
$O(\cdot)$	computational complexity
$\Gamma(\cdot)$	Gamma function
$R^n[\cdot]$	$n$ th order rank operator
$\text{Re}[\cdot]$	real part
$\text{Im}[\cdot]$	imaginary part
$E[\cdot]$	expectation
$ \cdot $	(element-wise) absolute value
$\ \cdot\ _p$	vector $L_p$ norm (default $L_2$ )



# Contents

<b>Abstract</b>	i
<b>Zusammenfassung</b>	iii
<b>List of Abbreviations</b>	v
<b>List of Symbols</b>	vii
<b>1 Introduction</b>	1
1.1 Motivations and Objectives	1
1.2 Thesis Outline	2
<b>2 Basics</b>	3
2.1 Single-look SAR Signal	3
2.1.1 <i>Point Scatterer</i>	3
2.1.2 <i>Distributed Scatterer</i>	3
2.1.3 <i>Layover of Scatterers</i>	4
2.2 Multipass InSAR	4
2.2.1 <i>Single-look InSAR Statistics</i>	5
2.3 The Relativeness of Multipass InSAR Estimates	6
2.4 Robust Estimation	8
2.4.1 <i>Maximum Likelihood Estimator (MLE)</i>	9
2.4.2 <i>M-estimator</i>	9
2.4.3 <i>Influence Function</i>	10
<b>3 State-of-the-art of Multipass SAR Interferometry</b>	12
3.1 State-of-the-art	12
3.1.1 <i>Persistent Scatterer Interferometry</i>	12
3.1.2 <i>Small Baseline Subset</i>	15
3.1.3 <i>SqueeSAR (Adaptive Multilooking)</i>	17
3.1.4 <i>SAR Tomography and Differential SAR Tomography</i>	18
3.1.5 <i>InSAR Point Cloud Fusion</i>	20
3.1.6 <i>Optical and SAR Images Fusion</i>	21
3.2 Contribution of this Thesis	22
3.2.1 <i>Computationally Efficient TomoSAR Inversion</i>	22
3.2.2 <i>TomoSAR Point Cloud Fusion</i>	22
3.2.3 <i>InSAR Optical Fusion</i>	22
3.2.4 <i>Robust Estimators</i>	22
<b>4 Summary of the Work</b>	23
4.1 Computationally Efficient Tomographic SAR Inversion	23
4.1.1 <i>Pre-classification of Single and Double Scatterers</i>	23
4.1.2 <i>Computational Cost Analysis</i>	26
4.1.3 <i>Tomographic Reconstruction of Entire City</i>	26
4.2 Automatic Feature-based TomoSAR Point Cloud Fusion	27
4.2.1 <i>Point Density Estimation</i>	28
4.2.2 <i>L-shape End Point Estimation</i>	29
4.2.3 <i>Fusion Result</i>	31
4.3 Semantic Interpretation of InSAR Estimates Using Optical Images	32
4.3.1 <i>Geometric Co-registration of Optical Images and InSAR Point Cloud</i>	33

4.3.2	<i>TomoSAR Point Cloud Rendering</i>	34
4.3.3	<i>Practical Demonstration</i>	39
4.4	<b>Robust Estimators for Multipass SAR Interferometry</b>	42
4.4.1	<i>Robust Covariance Matrix Estimators</i>	42
4.4.2	<i>Generalized Rank Covariance Estimator</i>	47
4.4.3	<i>Robust Phase History Parameters Estimators</i>	49
4.4.4	<i>Practical Demonstration of the Robust InSAR Optimization</i>	52
<b>5</b>	<b>Conclusion and Outlook</b>	58
5.1	Conclusion	58
5.2	Outlook	59
	<b>References</b>	61
	<b>Acknowledgement</b>	73
	<b>Appendices</b>	74
<b>A</b>	<b>Wang Y, Zhu X, Bamler R (2014) An Efficient Tomographic Inversion Approach for Urban Mapping Using Meter Resolution SAR Image Stacks. IEEE Geoscience and Remote Sensing Letters, 11 (7): 1250 - 1254.</b>	75
<b>B</b>	<b>Wang Y, Zhu X (2015a) Automatic Feature-based Geometric Fusion of Multi-view TomoSAR Point Clouds in Urban Area. IEEE Journal of Selected Topics in Applied Earth Observation and Remote Sensing, 8 (3): 953 - 965.</b>	83
<b>C</b>	<b>Wang Y, Zhu X, Zeisl B, Pollefeys M (2015b) Fusing Meter-Resolution 4-D InSAR Point Clouds and Optical Images for Semantic Urban Infrastructure Monitoring. IEEE Transactions on Geoscience and Remote Sensing. submitted.</b>	99
<b>D</b>	<b>Wang Y, Zhu X, Bamler R (2012) Retrieval of Phase History Parameters from Distributed Scatterers in Urban Areas Using Very High Resolution SAR Data. ISPRS Journal of Photogrammetry and Remote Sensing, 73: 89 - 99.</b>	143
<b>E</b>	<b>Wang Y, Zhu X (2015c) Robust Estimators in Multipass SAR Interferometry. IEEE Transactions on Geoscience and Remote Sensing, 54 (2): 1 - 13.</b>	157
<b>F</b>	<b>Publishers Copyright Policies</b>	173

# 1 Introduction

## 1.1 Motivations and Objectives

**S**ynthetic aperture radar (SAR) is an important remote sensing means that emits electromagnetic wave to the imaged object and receives the backscattered echoes. SAR interferometry (InSAR) is the techniques of using SAR as interferometer for 3-D reconstruction, and especially for monitoring subwavelength level deformation. InSAR and InSAR algorithms play irreplaceable roles in earth observation, because it has been generally appreciated that spaceborne InSAR is so far the only method to assess millimeter-precision long-term ground deformation on a global scale.

The real beauty of spaceborne InSAR is the repeated long-term monitoring of small deformation over large areas. The corresponding techniques are known as multipass / multi-baseline / multitemporal InSAR techniques. They make use of a stack of co-registered SAR images to reconstruct the deformation history, as well as the 3-D position of the scatterer.

The main focus of this thesis is on the algorithm development of multipass InSAR techniques for future SAR systems.

The last several years are the era of meter-resolution spaceborne SAR sensors, such as TerraSAR-X (TS-X), TanDEM-X and COSMO-SkyMed. That opened a whole new perspective of analysing spaceborne SAR data. Substantial development in multipass InSAR techniques has been made, e.g. in monitoring individual buildings, modelling higher order motion, reconstruction of 3-D objects from SAR image stacks, and so on.

In the next several years, SAR remote sensing will still be one of the most important means of spaceborne earth observation. In particular, more and more high resolution wide swath missions are under development. For example, the L-band SAR mission Tandem-L will provide one meter spatial resolution with a swath of 50km in its high resolution mode; the ALOS-2 provides both meter-resolution images with swath of 25km and 50km; and the TS-X Next Generation that will have decimetre-level spatial resolution. For the medium resolution mission, we can also foresee a data rush of the SAR satellite Sentinel-1, because of the free data access policy.

It is obvious that the future earth observation will, on one hand, have huge demand on large area processing, and on the other hand face data with unprecedented high resolution. The ever increasing higher resolution brought challenges, e.g. exponential growth of data volume, redundancy of data, different observation statistics, and much more dynamic motion. The challenges posed to my research under the context of multipass InSAR algorithms can be summarized as 1) How to efficiently perform tomographic SAR inversion for large data, because it is the most competent means of 3-D or even 4-D SAR imaging, but also the most computationally expensive one. 2) How to fuse meter-resolution SAR tomographic point clouds of urban areas from multiple viewing angles? One must consider the much higher point density compared to other InSAR methods, e.g. from persistent scatterer interferometry. 3) how to make use of the complementary information in high resolution optical image to help the interpretation of InSAR point clouds of urban areas, as an unstructured InSAR point cloud only gives 3-D position, deformation parameters, and the reflectivity of the scatterers, and lastly 4) how to handle unexpected statistical properties for unprecedentedly high resolution SAR images, e.g. the covariance matrix estimation of non-Gaussian scatterers.

Therefore, four important objectives of the thesis have been identified:

- ◇ Computational efficiency of SAR tomographic inversion: develop computationally efficient tomographic SAR inversion algorithms for very high resolution urban monitoring. The algorithm should be able to deliver tomographic reconstruction of an entire city using meter-resolution data in an affordable time.
- ◇ Information fusion: fuse TomoSAR point clouds from multiple viewing angles, in order to have a complete point cloud of the city under monitoring. The algorithm should be able to extract important feature points from the dense TomoSAR point cloud, in order to reduce computational cost and to be robust against large amount of outliers contained in the data, e.g. caused by the complex scattering scenario (Auer et al., 2011).
- ◇ Contextual awareness: fuse optical images and InSAR point cloud/SAR images for better interpretation of SAR images and the deformation signal. The algorithm should particularly address the problem of 3-D geometric registration of optical and SAR images, because of the complex 3-D urban topography.
- ◇ Statistical robustness: develop statistically robust InSAR algorithms for non-urban area monitoring, taking into account different SAR statistics that may bias the covariance estimation and much more dynamic deformation signal which can lead to model underfitting.

## 1.2 Thesis Outline

This is a cumulative dissertation comprising of five full paper peer-reviewed journal papers (one still under review), which are presented in the appendix of the thesis. Chapter 2 covers the necessary basics for the understanding of the thesis. The contribution of the thesis and the state-of-the-art are summarized in Chapter 3. Chapter 4 and Chapter 5 cover the summary of the five journal papers and the conclusion, respectively. Therefore, advanced readers can directly start from Chapter 4.

The variables in this thesis were written with the following rule: *italic*: scalar; bold regular: vector, and capital bold regular: matrix. The list of symbols can be found in List of Symbols.

## 2 Basics

Targeting at the readers who are already in the research field of InSAR, the necessary basics are included in this chapter. The discussions in this chapter are based on co-registered complex-valued SAR image stacks. For the basics of SAR imaging geometry, acquisition, focusing, etc., the readers are kindly directed to (Bamler and Hartl, 1998; Rosen et al., 2000) for an extensive review of these fundamentals.

### 2.1 Single-look SAR Signal

“Single-look” SAR signal refers to the reflected echoes integrated into one range-azimuth resolution cell, i.e. a range-azimuth SAR pixel. It is also known as single look complex (SLC). And in the rest of this thesis, they will be used interchangeably, unless explicitly stated. For the mathematical convenience, the classical SAR and InSAR theory has restricted the discussion to two extreme cases of scattering object: distributed scatterer and point scatterer.

#### 2.1.1 Point Scatterer

Point scatterer refers to the situation when the signal reflected from a single object is absolutely dominating in the resolution cell. The complex radar return  $g$  is modelled as a deterministic signal which is (Bamler and Hartl, 1998)

$$g = A \exp\left(-j\frac{4\pi}{\lambda}R + j\varphi_{scatter}\right) \quad (2.1)$$

where  $A$  is the signal amplitude,  $-4\pi R/\lambda$  and  $\varphi_{scatter}$  are the phase contribution from the distance  $R$  to the sensor and an intrinsic scattering phase, respectively. If the reflectance from the point scatterer is coherent throughout the whole stack of SAR images, it is then known as persistent/permanent scatterer (PS). Therefore, the scattering phase  $\varphi_{scatter}$  of a PS is usually assumed to be identical throughout the stack.

#### 2.1.2 Distributed Scatterer

Distributed scatterer (DS) or Gaussian scatterer is the superposition of sufficiently many random sub-scatterers within a resolution cell. By assuming none of the sub-scatterer dominates another, the complex backscattering of the resolution cell degenerates into a zero-mean complex circular Gaussian (CCG) random variable according to so-called *central limit theorem*, i.e.:

$$g = x + jy \quad x, y \sim N(0, \sigma^2) \quad (2.2)$$

where the real and imaginary part  $x$  and  $y$  both follow a zero-mean Gaussian distribution with variance  $\sigma^2$ . This assumption is fulfilled for natural objects in medium resolution ( $\sim 10$  m) SAR system, however can be violated in urban areas or for high resolution systems.

As both the real and imaginary parts are identically Gaussian distributed, one can derive that the intensity of a DS pixel  $I = |g|^2$  follows a  $\chi^2$  distribution with two degrees of freedom (DoF). The joint probability density function (PDF) of its intensity and phase is again a  $\chi^2$  distribution but only scaled with a factor of  $1/2\pi$  (Bamler and Hartl, 1998):

$$f(I, \varphi) = \frac{1}{2\pi I} \exp\left(-\frac{I}{\bar{I}}\right) \quad (2.3)$$

where  $\bar{I} = \sigma^2$  is the expected intensity. As one can see, the PDF does not depend on the phase, because the scattering phase  $\varphi_{scatter}$  of single-look DS has a uniform distribution. As a consequence, the phase seemingly carries no information.

### 2.1.3 Layover of Scatterers

Layover of scatterers happens when multiple scatterers appear at different elevation in the same resolution cell. The complex single look signal is then the integration of all the contributions along the third SAR coordinate elevation which is orthogonal to range and azimuth. Recall the plane wave approximation of the SAR imaging model of a single-look SAR pixel (after focusing) (Bamler and Hartl, 1998; Zhu, 2008):

$$g = \int_{\Delta s} \gamma(s) \exp\left(-j \frac{4\pi}{\lambda} R(s)\right) ds \quad (2.4)$$

where  $\gamma(s)$  is the complex reflectivity profile along the elevation direction  $s$ ,  $\Delta s$  is the extend of the elevation direction, and  $R(s)$  is the range change along elevation direction. As one can see, an SLC SAR image is only a single tomographic projection of the imaged object/area. It requires multiple acquisitions from slightly different positions in order to reconstruct the full reflectivity profile  $\gamma(s)$ . This is known as *SAR tomography* (TomoSAR).

In urban areas, we mostly consider the layover of PSs in a resolution cell. Therefore,  $\gamma(s)$  can be parametrically modelled as multiple delta functions:

$$\gamma(s)_{PS} = \sum_i A_i \delta(s - s_i) \quad (2.5)$$

where  $A_i$  and  $s_i$  are the unknown complex amplitude and the elevation position of the  $i$ th scatterer, respectively.

## 2.2 Multipass InSAR

Briefly speaking, InSAR is the technique of using SAR as interferometer. Figure 1 shows the plane wave approximation of the interference of two SAR images. The red and blue lines are the iso-range lines with interval of half wavelength of the two images respectively. Because of the perpendicular baseline  $B_\perp$ , the iso-range lines intersect each other, creating an interference pattern. The background fringe shows the interferometric phase of the pattern. The topography is mapped into interferometric phase depending on the height of the point.

Since the perpendicular baseline is much smaller than the range of the satellite to the imaged object, every iso-range line defined by  $R = \text{constant}$  in the two SAR images should cover the same range resolution cell. This is a prerequisite of the coherent nature of SAR imaging, i.e. the two images should observe the same object in their corresponding pixels.

The interferometric phase changes as the baseline  $B_\perp$  changes. The unambiguous interferometric phase  $\phi$  is shown to be linearly proportional to the topography height  $h$  with the equation:

$$\phi = -\frac{4\pi}{\lambda} \frac{B_\perp}{R_S \sin \theta} h + \phi_{defo} \quad (2.6)$$

where  $\lambda$  is the wavelength,  $R_S$  is the nominal range which is the range of the SAR sensor to a zero-elevation point, and  $\theta$  is the incidence angle. The deformation phase  $\phi_{defo}$  must be taken into account if the two images are acquired at different time. The deformation causes

absolute phase change which is not dependent on the baseline. If the pixel is a DS, then (2.6) is the expected interferometric phase.

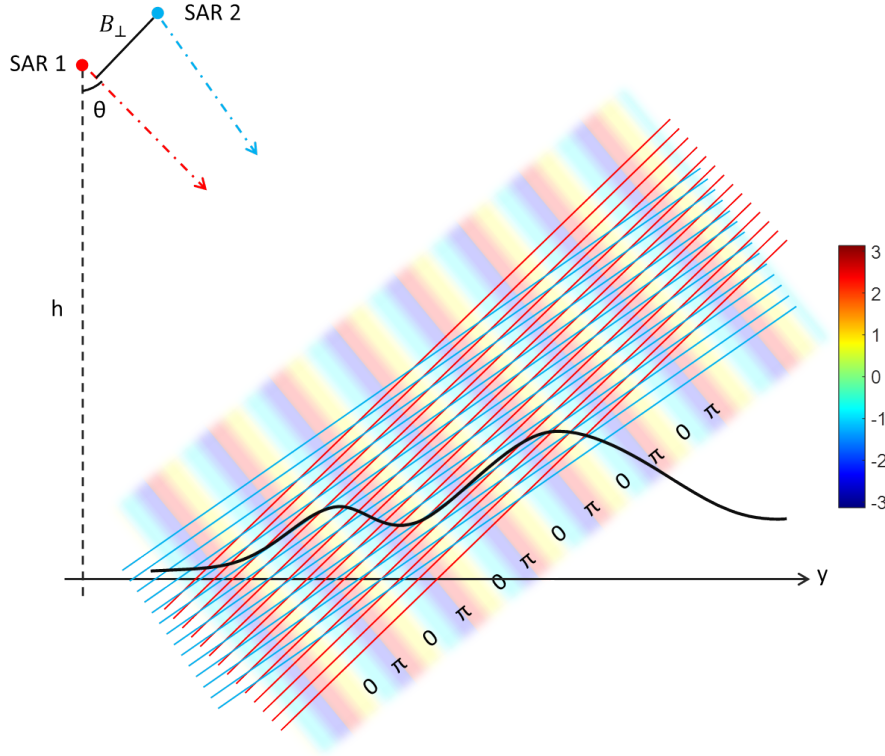


Fig. 2.1. Plane wave approximation of the interference of two SAR images. The red and blue lines are the iso-range lines with interval of half wavelength of the two images respectively. The iso-range lines intersect each other, creating an interference pattern. The background shows the interferometric phase of the pattern. The topography is mapped into interferometric phase depends on the height of the point.

Multipass InSAR is an extension of the single-baseline InSAR. It exploits the phase difference of more than two complex-valued SAR images. It is often employed for long-term deformation monitoring, which is also known as differential InSAR. One can derive dynamic geophysical quantities such as deformations rate (volcanoes, urban underground water, oil reservoir, etc.).

As long as only the phase is concerned, the unambiguous interferometric phases  $\Phi$  of the multiple acquisitions with their perpendicular baselines  $\mathbf{B}_\perp$  are:

$$\Phi = -\frac{4\pi}{\lambda} \frac{\mathbf{B}}{R_S \sin \theta} h + \Phi_{defo}. \quad (2.7)$$

The deformation phase vector  $\Phi_{defo}$  is often modelled as a function  $d(\mathbf{t}_n)$  (e.g. linear or periodic) of the acquisition time  $\mathbf{t}_n$ . The details of popular multipass InSAR techniques are covered in Chapter 3.

### 2.2.1 Single-look InSAR Statistics

It has been mentioned before that equations (2.6) and (2.7) only apply to PS. Under the assumption of distributed scatterers, they represent the expected interferometric phase. The phase variance caused by the decorrelation between the two co-registered SAR pixels  $g_1$  and  $g_2$  is fully characterized by their complex covariance. Normalizing the covariance with the expected intensities of the pixels, we obtain a more commonly used quantity - the complex

coherence  $\gamma$ :

$$\gamma = \frac{E[g_1 g_2^*]}{\sqrt{E[|g_1|^2] E[|g_2|^2]}}. \quad (2.8)$$

The PDF of the interferometric sample  $v = g_1 g_2^*$ , represented by the joint PDF of its amplitude and the interferometric phase can be shown as follows (Lee et al., 1994; Tough et al., 1995; Bamler and Hartl, 1998)

$$f_{|v|, \phi}(|v|, \phi) = \frac{2|v|}{\pi \bar{I}^2 (1 - |\gamma|^2)} \exp\left(\frac{2|\gamma||v| \cos(\phi - \bar{\phi})}{\bar{I}(1 - |\gamma|^2)}\right) K_0\left(\frac{2|v|}{\bar{I}(1 - |\gamma|^2)}\right) \quad (2.9)$$

where  $\bar{I} = \sqrt{E[|g_1|^2] E[|g_2|^2]}$ ,  $\bar{\phi}$  is the expected interferometric phase, and  $K_0$  is the modified Bessel function of the second kind. Generalizing to multibaseline SAR pixels, the joint PDF of the complex vector  $\mathbf{g}$  is a multivariate CCG random variable. The joint PDF can be written as the following compact equation:

$$f_{\mathbf{g}}(\mathbf{g}) = \frac{1}{\pi^N \det(\mathbf{C})} \exp(-\mathbf{g}^H \mathbf{C}^{-1} \mathbf{g}) \quad (2.10)$$

where  $N$  is the number of images, and  $\mathbf{C}$  is the complex covariance matrix whose analytical expression is as follows.

$$\mathbf{C} = \begin{bmatrix} \bar{I}_1 & \gamma_{1,2} \sqrt{\bar{I}_1 \bar{I}_2} & \cdots & \gamma_{1,N} \sqrt{\bar{I}_1 \bar{I}_N} \\ \gamma_{1,2}^* \sqrt{\bar{I}_1 \bar{I}_2} & \bar{I}_2 & & \\ \vdots & & \ddots & \vdots \\ \gamma_{1,N}^* \sqrt{\bar{I}_1 \bar{I}_N} & & \cdots & \bar{I}_N \end{bmatrix} \quad (2.11)$$

As one can see in the CCG case, the multivariate  $\mathbf{g}$  is fully characterized by the complex covariance matrix  $\mathbf{C}$ . Therefore the estimation of the covariance matrix becomes an important step in many multipass InSAR techniques. Under the assumption CCG distribution, the maximum likelihood estimator (MLE) of the covariance matrix can be easily derived given a set of  $M$  spatially stationary samples  $\mathbf{G} = [\mathbf{g}_1, \mathbf{g}_2, \dots, \mathbf{g}_M]$ :

$$\hat{\mathbf{C}}_{MLE} = \frac{1}{M} \sum_{m=1}^M \mathbf{g}_m \mathbf{g}_m^H = \frac{1}{M} \mathbf{G} \mathbf{G}^H \quad (2.12)$$

## 2.3 The Relativeness of Multipass InSAR Estimates

The InSAR estimates, i.e. the topography and deformation estimates, are relative with respect to a reference point in the image that was selected during the processing. The unknown topography and deformation of the reference point shift the estimates of the whole image along the elevation direction of the native radar coordinate system. When the InSAR estimates are transformed to geographical coordinate systems, it will eventually cause the position to shift in all the three dimensions, in addition with a scaling effect due to the different incidence angle at different range.



The following figure explains the aforementioned systematic shift in the Universal Transverse Mercator (UTM) coordinate system. The two black dots indicate the true positions of two scatterers which are on the same height w.r.t. a reference surface. The two scatterers are located at the near range and far range of the SAR image. Their incidence angles  $\theta_{near}$  and  $\theta_{far}$  are different, e.g. as many as one degree for TS-X spotlight image. Of course, the difference is exaggerated in the plot for a better visualization. Due to the height  $\Delta h$  of the reference point, the scatterers are shifted to the red points along their elevation directions, respectively. As one can see that the scatterers are equally shifted in height, but scaled in the ground range direction due to their different elevation directions. The difference between the scaled ground range and the true value is

$$\Delta R = \frac{\Delta h}{\tan \theta_{near}} - \frac{\Delta h}{\tan(\theta_{near} + \Delta \theta)} \quad (2.13)$$

$$\approx \Delta h \Delta \theta \quad \text{if } \theta_{near} \gg \Delta \theta$$

where  $\theta_{far} = \theta_{near} + \Delta \theta$ . Such difference is approximately linearly proportional to the unknown height of the reference point and the diversity of the incidence angle. As the heading angle of the satellite is not parallel to north-south direction, this scaling will also be transferred to both north-south and east-west directions.

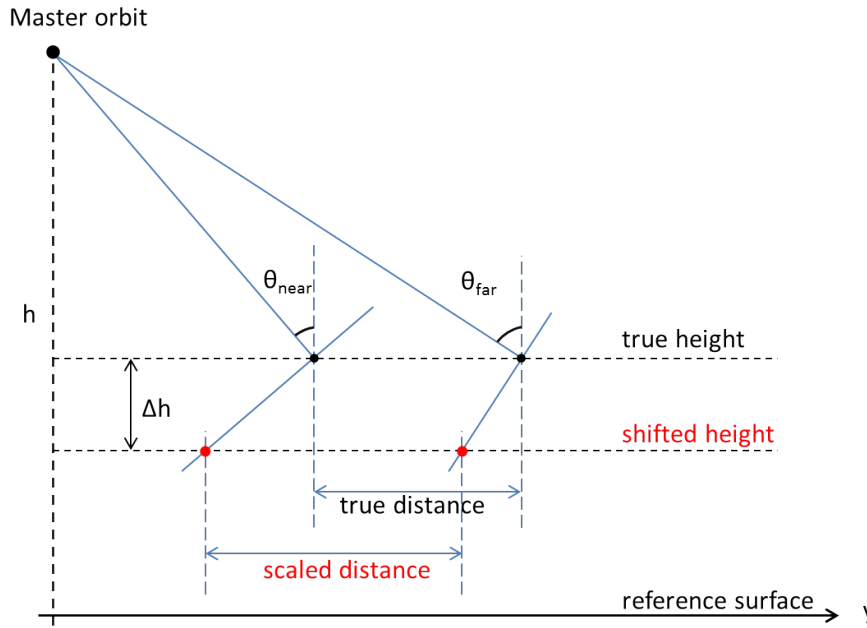


Fig. 2.2. Illustration of systematic shift of the multipass InSAR position estimates in UTM coordinate. The two black dots indicate the true positions of two scatterers which are on the same height w.r.t. a reference surface. The two scatterers are located at the near range and far range of the SAR image, respectively. Their incidence angles  $\theta_{near}$  and  $\theta_{far}$  are different, e.g. as many as one degree for TS-X spotlight images. Due to the unknown height of the reference point, the scatterers are shifted to the red points along their elevation directions, respectively. The scatterers are equally shifted in height, but scaled in the ground range direction due to their different elevation directions. This ground range scaling will translate into both north and east direction in the UTM coordinate. Therefore, any relative InSAR position estimate is shifted in three directions, with additional horizontal 2-D scaling.

Figure 2.3 also plots the ground range difference per meter of the height of the reference point as a function of the near range incidence angle and the angle diversity. As one can see, the approximation in (2.13) is only applicable for large incidence angles. For a typical image stack of  $40^\circ$  of incidence angle and  $1^\circ$  diversity, the scaling is 3.5 cm/m/°. However, the scaling will become 15 cm/m/° if the incidence angle becomes  $20^\circ$ , which will cause 1.5m of scaling if the unknown height of the reference point is 10 m.

The shift can be mitigated by choosing a reference point close to the reference digital elevation model (DEM), i.e.  $h \approx 0$ m. For medium resolution SAR data of non-urban area, such

shift is not easily noticeable, because of the relatively smooth topography and large pixel size. However, they must be taken into account for meter resolution SAR data of urban area.

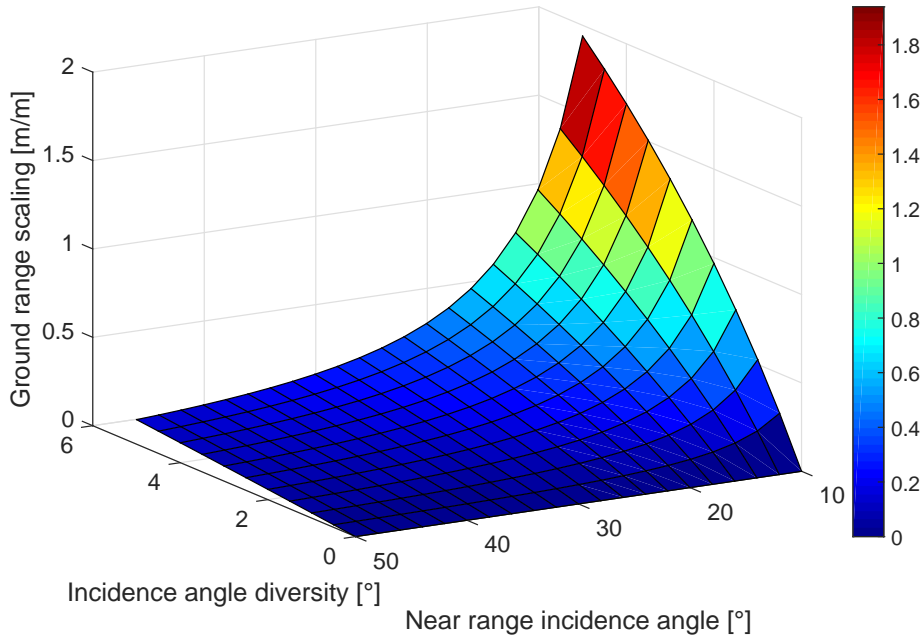


Fig. 2.3. Maximum ground range scaling (per meter of the height of reference point) as a function of the near range incidence angle and the incidence angle diversity between near range and far range. For a typical image stack of  $40^\circ$  incidence angle and  $1^\circ$  diversity, the scaling is  $3.5 \text{ cm/m}^\circ$ . However, the scaling will become  $15 \text{ cm/m}^\circ$  if the incidence angle becomes  $20^\circ$ , which will cause 1.5 m of scaling if the unknown height of the reference point is 10 m.

An more accurate method to mitigate this shift is to find the absolute 3-D position of the reference point. This concept is the so-called *geodetic InSAR* in which: by means of "SAR imaging geodesy" (Eineder et al., 2011; Cong et al., 2012; Balss et al., 2014) the absolute 2-D (range and azimuth) positions of targets such as corner reflectors and persistent scatterers can be estimated to centimeter-level accuracy after compensating for the most prominent geodynamic and atmospheric error sources; utilizing the stereo configuration available from multiple viewing angles, the absolute 3-D positions of the selected points can be retrieved using StereoSAR (Gisinger et al., 2015); finally, the InSAR estimates are reconstructed with respect to these points whose absolute positions are known. In this manner, the obtained geodetic InSAR estimates are absolute (Zhu et al., 2015c). However, this strategy relies on the successful identification of exactly identical scattering object in SAR images acquired from different view angles which is very challenging (Gisinger et al., 2015; Zhu et al., 2015c).

## 2.4 Robust Estimation

The estimator of a parameter is derived based on certain assumptions. Given the PDF of the measurement noise, the MLE is the best unbiased estimator if no prior knowledge of the parameter is given. In practice, we should also consider the robustness of the estimator. A robust estimator should degrade gracefully with respect to the violation of the assumption. For multipass InSAR techniques, we are interested in the performance of the estimator with respect to different SAR signal and phase statistics.

The performance of an estimator is reflected in the variance and bias of its estimate. An estimator which is unbiased and attains the Cramér-Rao lower bound (CRLB) is said to be *efficient* (Kay, 2001). Therefore, the term *efficiency* in the context of estimator often refers the ratio of the estimator's variance w.r.t. the CRLB.

In the following, the basics of MLE will be recapped in order to introduce the M-estimator which brings a unified view of weighted averaging.

### 2.4.1 Maximum Likelihood Estimator (MLE)

An MLE maximizes the goodness-of-fit (likelihood) of the measurements to the parameters to be estimated. It is always with respect to a certain distribution of the measurement statistics. Consider a simple location estimation problem, where the likelihood function of the univariate observation of the location  $g \in \mathbb{R}$  is  $f_g(g|\mu)$  in which  $\mu$  is the expected location to be estimated. Let  $f_g(g|\mu)$  equal a known PDF  $f_x(x)$ , where  $x = g - \mu$ . Without losing generality, let us assume that  $x$  is scaled to unit variance. Given multiple independent observations of  $g$ , the MLE of  $\mu$  maximizes the product of the likelihood:

$$\hat{\mu}_{\text{MLE}} = \arg \max_{\mu} \prod_i f_x(x_i) = \arg \min_{\mu} \sum_i -\ln f_x(x_i) \quad (2.14)$$

The above equation is usually done in the logarithmic domain, because  $f_x(x)$  is often in certain exponential form, e.g. Gaussian distribution under which (2.14) is simplified to

$$\hat{\mu}_{\text{MLE}} = \arg \min_{\mu} \sum_i (g_i - \mu)^2 = \text{average}(g_i) \quad (2.15)$$

### 2.4.2 M-estimator

M-estimators are a class of well-known robust estimators that take the form of a minimization problem:  $\min \sum_i \rho(x_i)$ . The M-estimator permits a minimization of a customized loss function  $\rho(x)$  of the residuals instead of only the quadratic form, to resist outliers. The M-estimator of  $\mu$  is as follows:

$$\hat{\mu}_{\text{M-est}} = \arg \min_{\mu} \sum_i \rho(x_i) \quad (2.16)$$

Therefore, choosing  $\rho(x) = -\ln f_x(x)$  is equivalent to the MLE. For example, the MLE of Gaussian distributed residual is an M-estimator with  $\rho(x) = x^2$ .

Equation (2.16) can be solved by finding the zeros of its derivative, which can be written as

$$\sum_i \frac{\rho'(x_i)}{x_i} x_i = 0 \quad (2.17)$$

where  $\rho'(x)$  is the derivative of  $\rho(x)$ .  $\rho'(x)/x$  is commonly referred to as the weighting function  $w(x)$ , because substituting  $x_i = g_i - \mu$  into the previous equation yields an M-estimator that is actually a weighted average of the observations:

$$\hat{\mu}_{\text{M-est}} = \frac{\sum_i w(g_i - \hat{\mu}) g_i}{\sum_i w(g_i - \hat{\mu})} \quad (2.18)$$

Equation (2.18) can be solved iteratively with an initial estimate  $\hat{\mu}$  being given and the weights being updated at each iteration. Therefore, an M-estimator is actually an iteratively re-weighted averaging, and the weighting function can be directly derived from the PDF

$f_x(x)$  (if exist) of the observations with the following relation

$$w(x) = \frac{-\partial \ln f_x(x)}{x \partial x} \quad (2.19)$$

Typically, the weighting function is designed to have high efficiency for normal distribution. Therefore, a more common form of the weights is  $w = \rho'(x/\sigma)/(x/\sigma)$  where  $\sigma$  is the standard deviation of  $x$ . Should the PDF be unknown, one may use empirical robust weighting function, such as the Tukey biweight functions, such as the Tukey biweight function (Tukey, 1977). The convergence of (2.18) is only guaranteed for convex  $\rho$  functions (Venables and Ripley, 2002). The initial estimate must be carefully estimated should  $\rho$  be nonconvex.

To summarize, M-estimator is a trade-off between robustness and efficiency. Nevertheless, it can still maintain satisfactory efficiency under the nominal model by properly choosing the loss function.

### 2.4.3 Influence Function

The influence function qualitatively measures the robustness of the estimator. It describes the influence of infinitesimal outlier at an arbitrary point on the estimator. Given the nominal distribution  $F$  of the observation, the fraction of contamination  $\varepsilon$ , and the estimate  $\hat{g}$ , the influence function is defined as follows:

$$IF(x|\hat{g}, F) = \lim_{\varepsilon \rightarrow 0^+} \frac{\hat{g}(G) - \hat{g}(F)}{\varepsilon} \quad (2.20)$$

where  $G = \varepsilon \Delta_x + (1 - \varepsilon)F$  is the contaminated distribution with  $\Delta_x$  being the point-mass probability.

The influence function of a qualitatively robust estimator should be bounded and continuous, which means a small fraction of outliers can have only limited effect on the estimates, and gradual changes in the data also lead to gradual changes in the estimate (Zoubir et al., 2012).

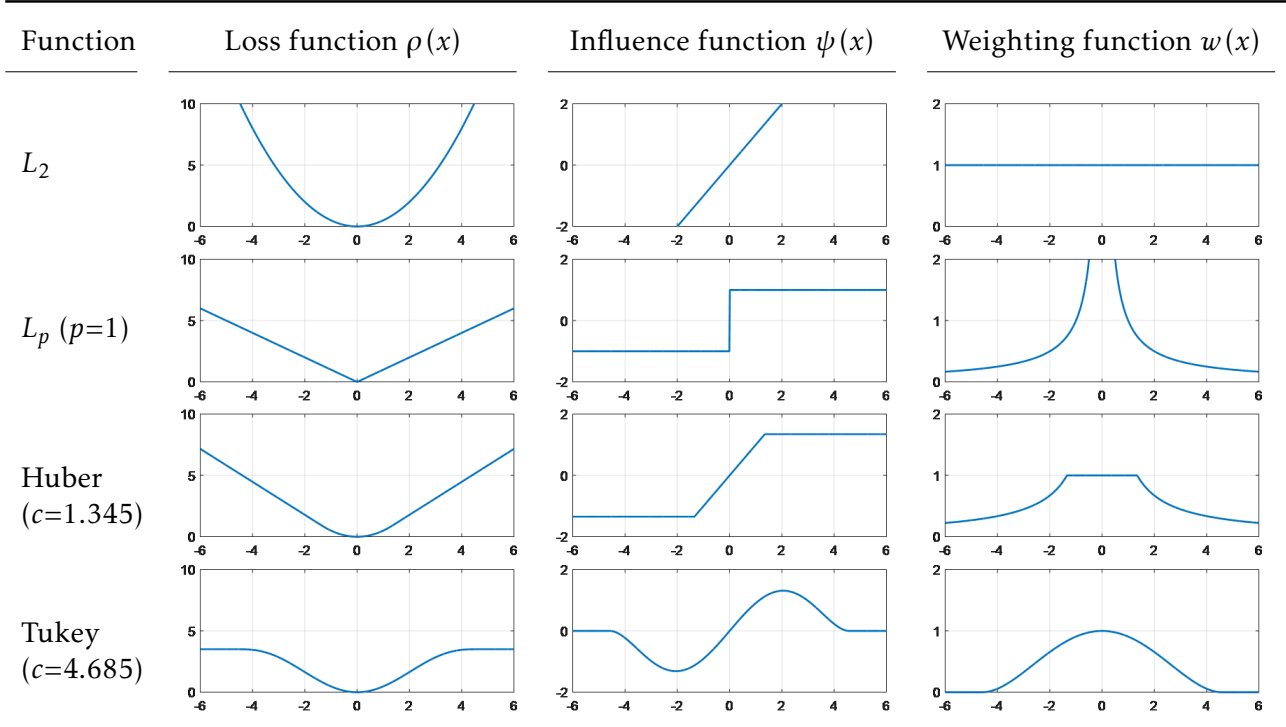
For M-estimators, the influence function with nominal distribution being Gaussian is proportional to the derivative of the loss function, i.e.  $IF \propto \psi(x) = \rho'(x)$ . This allows us to conveniently study the performance of different robust loss functions. Table 1 shows some typical loss functions, and the corresponding influence and weighting functions. And their plots are also shown in Table 2.

The  $L_2$  loss function corresponds to Gaussian distributed noise. Its influence function is not bounded. One large outlier will heavily bias the estimates. If we generalize to  $L_p$  norm minimization, e.g.  $L_1$ , its influence function is bounded, meaning the influence of large outliers is limited. Another popular robust loss function is the Huber function named after P.J. Huber — the pioneer in robust estimation (Huber, 1981). Tukey biweight function (Tukey, 1977) is one of the re-descending functions, since its influence function is bounded and descends to zeros as the position of the outlier increases to a certain value. The parameter  $c \in \mathbb{R}^+$  in Huber and Tukey functions is to tune the performance of the estimator. The lower the value of  $c$ , the more robust but less efficient is the estimator for Gaussian distributed noise, and vice versa. Therefore,  $c$  is usually chosen to give a relative efficiency of 95% at normal distribution. This number is 1.345 and 4.586 for Huber and Tukey functions, respectively (Venables and Ripley, 2002).

Table 1. Some typical loss functions, and their corresponding influence and weighting functions.

Function	Loss function $\rho(x)$	Influence function $\psi(x)$	Weighting function $w(x)$
$L_2$	$x^2/2$	$x$	1
$L_p$	$ x ^p/p$	$x x ^{p-2}$	$ x ^{p-2}$
Huber	$\begin{cases}  x  < c & x^2/2 \\  x  \geq c & c x  - c^2/2 \end{cases}$	$\begin{cases}  x  < c & x \\  x  \geq c & c \operatorname{sign}(x) \end{cases}$	$\begin{cases}  x  < c & 1 \\  x  \geq c & c/ x  \end{cases}$
Tukey	$\begin{cases}  x  < c & c^2/6 - (c^2 - x^2)^3/6c^4 \\  x  \geq c & c^2/6 \end{cases}$	$\begin{cases}  x  < c & x(c^2 - x^2)^2/c^4 \\  x  \geq c & 0 \end{cases}$	$\begin{cases}  x  < c & (c^2 - x^2)^2/c^4 \\  x  \geq c & 0 \end{cases}$

Table 2. Plots of the loss functions, and their corresponding influence and weighting functions.



### 3 State-of-the-art of Multipass SAR Interferometry

This chapter will focus on the estimation aspect of the state-of-the-art multipass InSAR techniques including persistent scatterer interferometry (PSI), small baseline subset (SBAS), SqueeSAR, and TomoSAR.

The discussion is restricted to the two extreme cases of scattering objects: PS and DS. DSs are the most natural scatterers in a SAR image. Because of its scattering mechanism, it suffers from temporal and geometric decorrelation. To reduce the effect of decorrelation, one can either use interferograms of small spatial and temporal baselines, i.e. SBAS, or average pixels of same scattering statistics, i.e. SqueeSAR.

PS overcomes the limitation of DS by identifying single scatterers whose phase is stable throughout the SAR image stack. Theoretically, PSs do not suffer from geometrical decorrelation, permitting the use of interferograms with large spatial baseline even beyond the critical one.

TomoSAR is a generalization of the single phase center PS or DS model, by considering the full reflectivity profile along the radar third dimension elevation. It solves the layover of scatterers in a range-azimuth resolution cell.

#### 3.1 State-of-the-art

##### 3.1.1 Persistent Scatterer Interferometry

###### General Remark

PSI was developed for long-term ground deformation monitoring (Ferretti et al., 2001). Due to the long temporal baseline, low resolution, and large spatial baseline of some SAR image stacks, the individual DS pixel often lose their coherence, rendering them not directly useful for long-term deformation monitoring. PSI was invented to exploit the PSs whose scattering phases are stable throughout the whole SAR image stack. Not only are the phases of PSs stable, but also is the dimension of a PS often much smaller than a resolution cell, permitting the use of image with large spatial baseline even beyond the critical one.

There exist several implementations of PSI (Ferretti et al., 2001; Adam et al., 2003; Hooper et al., 2004; Kampes and Adam, 2006) etc., as well as improvements using different motion models (Colesanti et al., 2003; Gernhardt and Bamler, 2012), improvements using dense differential PS pairs (Wang et al., 2014; Costantini et al., 2014), and some variants of PSI that exploit partially coherent PSs (Wegmüller et al., 2010; Perissin and Wang, 2012) which follow the same spirit of SBAS that will be covered in the next section. Generally, PSI estimates the atmospheric phase, the topography, and the motion parameters of each PS. A generally recognized workflow is as follows (Ferretti et al., 2001):

- (1)  $N$  interferograms are formed from  $N+1$  co-registered SAR images. A master scene is carefully selected to maximize the overall coherence. A reference DEM is subtracted from all the interferograms to flatten the topographic phase.
- (2) An initial set of PS candidates is selected, which should represent the set of most stable PSs in the scene. A few methods have been proposed to fulfil this task:
  - ◊ using amplitude dispersion analysis for high signal-to-noise-ratio (SNR) PSs (Ferretti et al., 2001). This is mostly applicable to urban areas.

- ◇ using phase analysis for low SNR PSs (Hooper et al., 2004), which was demonstrated to outperform the previous method in non-urban area.
  - ◇ and alternatively using signal-to-clutter-ratio to approximate the phase variance (Adam et al., 2004).
- (3) The PS candidates are connected to form a reference network. The differential PS pairs, also known as *arcs*, are limited to be within the spatial correlation length of typical atmospheric phase signal, so that the atmospheric phase can be greatly mitigated through differentiation. This length was found to be 2-3 km in (Hanssen, 2006). After that, the differential phase history parameters  $\theta$ , i.e. the differential topography height and differential motion parameters, of the arcs are estimated.

The differential height estimates are integrated with respect to a reference point, so that their absolute phase contribution can be removed from the interferometric phase, leaving the phase with contributions only from the deformation, atmosphere, and noise. The atmospheric phase is then extracted by low-pass filtering in the spatial domain followed by high-pass in the temporal domain, assuming the atmospheric phase is highly correlated in the spatial domain and temporally nearly uncorrelated for typical temporal baselines of tens of days.

The atmospheric phase is interpolated over the whole scene, so that it can be subtracted from the interferograms.

- (4) More PSs are detected after the mitigation of the atmospheric phase. They are linked to the nearest point in the reference network, and their phase history parameters are estimated.
- (5) The positions of PSs are geocoded to geographical coordinate system for further analysis.

### Maximum Likelihood Estimator

On the estimation point of view, a PS pixel value is modelled as a deterministic complex value with additive CCG noise. For a series of co-registered PSs,  $\mathbf{g}$ , its noise is assumed to follow an independent and identical distribution (i.i.d.). Under such assumption, the MLE of  $\theta$  is (Rife and Boorstyn, 1974; Wang and Zhu, 2015b)

$$\hat{\theta}_{MLE} = \arg \max_{\theta} \left\{ (2\pi\sigma^2)^{-N} \exp \left( -\frac{1}{2\sigma^2} \|\mathbf{g} - \bar{\mathbf{g}}(\theta)\|_2^2 \right) \right\} \quad (3.1)$$

where  $\theta$  is the vector of phase history parameters comprised of topography height  $h$ , linear deformation rate  $v$ , etc.,  $\bar{\mathbf{g}}(\theta)$  is the modelled PS signal, and  $\sigma^2$  is the variance of the complex noise. Simplifying the above equation leads to the well-known formulation of periodogram or the so called "temporal coherence" (Rife and Boorstyn, 1974; Colesanti et al., 2003; De Maio et al., 2009):

$$\hat{\theta}_{MLE} = \arg \max_{\theta} \left\{ \left| \frac{1}{N} \sum_{n=1}^N g_n \exp(-j\varphi_n(\theta)) \right| \right\} \approx \arg \max_{\theta} \left\{ \left| \frac{1}{N} \sum_{n=1}^N \frac{g_n}{|g_n|} \exp(-j\varphi_n(\theta)) \right| \right\} \quad (3.2)$$

where  $\varphi_n(\theta)$  is the modelled phase of the PS in the  $n$ th image. Often, the amplitude of  $\mathbf{g}$  is dropped in the estimation (Ferretti et al., 2001), since it barely changes the estimates for high SNR PSs.

Equation (3.2) can also be understood as parameter estimation of a complex signal with single frequency in each dimension of  $\theta$ , since the modelled phase  $\varphi_n(\theta)$  is linearly proportional to  $\theta$ . It finds the best fit of  $\varphi_n(\theta)$  with given  $N$  wrapped phases of  $\mathbf{g}$ .

### Least Squares Ambiguity Decorrelation (LAMBDA) Estimator

Equation (3.2) can also be treated as a phase unwrapping problem with parametric phase model. (Kampes and Hanssen, 2004) solved it using the LAMBDA method which is originally developed for the ambiguity resolution of GPS signal.

The LAMBDA method for PS parameter estimation is formulated as  $\boldsymbol{\varphi} = \mathbf{A}\mathbf{a} + \mathbf{B}\boldsymbol{\theta} + \boldsymbol{\varepsilon}$ , i.e.

$$\begin{bmatrix} \varphi_1 \\ \varphi_1 \\ \vdots \\ \varphi_N \end{bmatrix} = \begin{bmatrix} 0 & 0 & \cdots & 0 \\ -2\pi & & & \\ & -2\pi & & \\ & & \ddots & \vdots \\ 0 & & \cdots & -2\pi \end{bmatrix} \begin{bmatrix} a_2 \\ \vdots \\ a_N \end{bmatrix} + \begin{bmatrix} \xi_1 & \zeta_1 & \cdots \\ \xi_2 & \zeta_2 & \\ \vdots & \vdots & \\ \xi_N & \zeta_N & \cdots \end{bmatrix} \begin{bmatrix} h \\ v \\ \vdots \end{bmatrix} + \boldsymbol{\varepsilon} \quad (3.3)$$

where  $\boldsymbol{\varphi} \in \mathbb{R}^N$  is the wrapped phase of  $\mathbf{g}$ ,  $\mathbf{A}$  is a diagonal matrix of  $-2\pi$  but discarding the first column (since the ambiguities are relative),  $\mathbf{a} \in \mathbb{Z}^{N-1}$  is the integer ambiguity, matrix  $\mathbf{B}$  is the model matrix of the phase containing the parameter-to-phase conversion factors ( $\xi$ ,  $\zeta$ , etc.), and  $\boldsymbol{\varepsilon}$  is the phase noise.

The LAMBDA estimator is an integer least square problem:

$$\hat{\boldsymbol{\theta}}_{LAMBDA} = \arg \min_{\mathbf{a}, \boldsymbol{\theta}} \|\boldsymbol{\varphi} - \mathbf{A}\mathbf{a} - \mathbf{B}\boldsymbol{\theta}\|_2^2 \quad \text{s.t. } \mathbf{a} \in \mathbb{Z}^{N-1} \quad (3.4)$$

Equation (3.4) assumes a Gaussian phase noise which is not true for PS with CCG noise, but can be valid for PS of high SNR.

Considering the integer constraint of  $\mathbf{a}$ , solving (3.4) is a three-step procedure in (Kampes and Hanssen, 2004).

- (1) The unconstrained estimates  $\hat{\mathbf{a}}$  and  $\hat{\boldsymbol{\theta}}$ , also known as float solution, are computed by unconstrained least square. Important to note that the linear system of equation (3.3) is underdetermined, i.e. more unknowns than observations. Regularization, such as augmenting the system with pseudo-observations, must be introduced in order to solve the linear equation system.
- (2) Given the float solution  $\hat{\mathbf{a}}$ , its integer-constrained solution is equation (3.5).

$$\check{\mathbf{a}} = \arg \min_{\mathbf{a}} \|\hat{\mathbf{a}} - \mathbf{a}\|_{\mathbf{C}_{\hat{\mathbf{a}}\hat{\mathbf{a}}}}^2 \quad \text{s.t. } \mathbf{a} \in \mathbb{Z}^{N-1} \quad (3.5)$$

If each element of  $\hat{\mathbf{a}}$  is uncorrelated, i.e. the covariance matrix  $\mathbf{C}_{\hat{\mathbf{a}}\hat{\mathbf{a}}}$  is diagonal, (3.5) is a simple rounding. Otherwise, the correlation among the elements of  $\hat{\mathbf{a}}$  should be mitigated by multiplying it with an integer matrix  $\mathbf{Z}$  that decorrelates it the most. In another word, given the covariance matrix  $\mathbf{C}_{\hat{\mathbf{a}}\hat{\mathbf{a}}}$  of  $\hat{\mathbf{a}}$ , the covariance matrix of  $\hat{\mathbf{z}} = \mathbf{Z}^T \hat{\mathbf{a}}$  which is  $\mathbf{C}_{\hat{\mathbf{z}}\hat{\mathbf{z}}} = \mathbf{Z}^T \mathbf{C}_{\hat{\mathbf{a}}\hat{\mathbf{a}}} \mathbf{Z}$  should be as diagonal as possible. In practice,  $\mathbf{Z}$  matrix is found by the LDL decomposition (Golub and Loan, 1996) of  $\mathbf{C}_{\hat{\mathbf{a}}\hat{\mathbf{a}}}$ . To this end, (3.5) is transformed to

$$\check{\mathbf{z}} = \arg \min_{\mathbf{z}} \|\hat{\mathbf{z}} - \mathbf{z}\|_2^2 \quad \text{s.t. } \mathbf{z} \in \mathbb{Z}^{N-1} \quad (3.6)$$

And the solution search space has been greatly reduced. The diagonalness of  $\mathbf{C}_{\hat{\mathbf{z}}\hat{\mathbf{z}}}$  is not always guaranteed. Therefore, a series of bounds for  $\mathbf{z}$  is computed, in order to further restrict the search space.

- (3) The final integer-constrained solution  $\check{\boldsymbol{\theta}}$  is computed using least square after subtracting the  $2\pi$  ambiguities.



## Summary

To conclude, the estimation of phase history parameters of PS is rather straightforward, a benefit brought by the simple point scatterer model. The estimation techniques include the MLE (periodogram) and LAMBDA.

The accuracy of the estimates primarily depends on the number of images  $N$  and the SNR of the PS. Typically, for PSs with relatively high SNR, e.g. 10dB, it requires around 30 images to achieve a millimeter precision on the yearly linear deformation rate (Ferretti et al., 2001).

Comparing MLE and LAMBDA, they produce nearly identical results as demonstrated in (Gernhardt et al., 2007). Nevertheless, LAMBDA explicitly solves the ambiguity problem, whereas the periodogram does not. But periodogram will outperform LAMBDA for low SNR PSs, since the assumption of Gaussian phase noise in LAMBDA is no longer valid for low SNR PSs. However, LAMBDA is more computationally efficient than MLE (periodogram) because the ambiguity solution space in LAMBDA is greatly reduced, and the phase parameters are solved analytically by least square. Still, PSI is the most computationally efficient one among all state-of-the-art multipass InSAR methods (even periodogram is used). Therefore, it is widely used, and has become the workhorse for deformation monitoring.

### 3.1.2 Small Baseline Subset

#### General Remark

SBAS exploits DS instead of PS. As we know, DSs will suffer from temporal decorrelation. The innovation of SBAS (Berardino et al., 2002) is to form interferograms of only small baselines, so that the geometric and temporal decorrelation of DSs can be controlled. The following figure shows the baseline plot of a single master InSAR stack (e.g. PSI) and SBAS. The scattered dots are perpendicular baseline versus temporal baseline of each image. PSI links every image to a single master, while SBAS only forms interferograms with small spatial and temporal baselines, resulting in multiple masters.

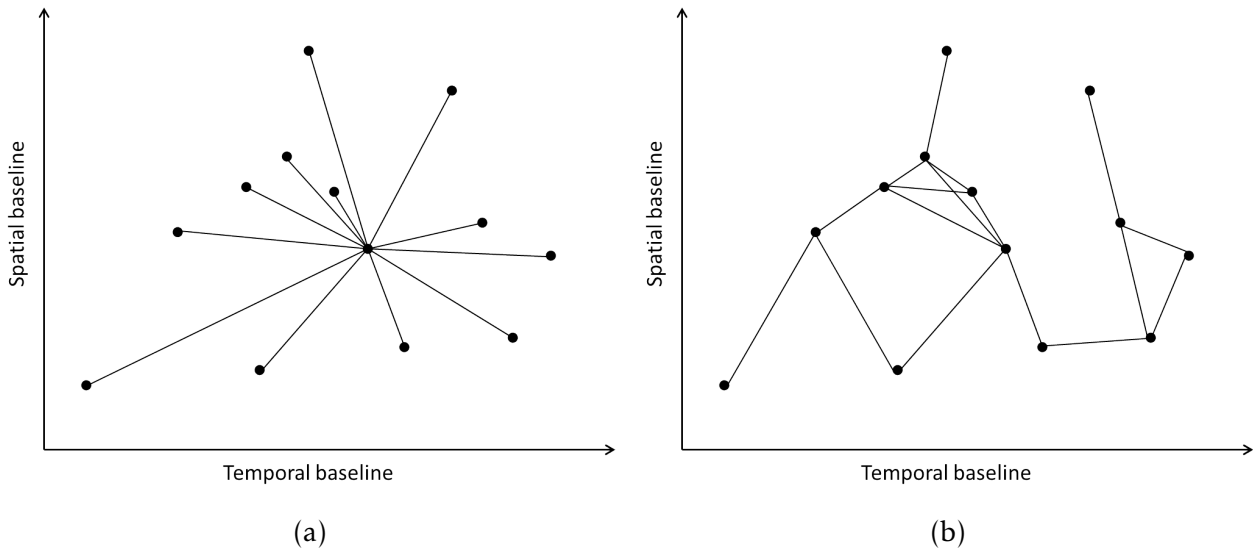


Fig. 3.1. Spatial-temporal baseline plot of (a) single master InSAR stack, e.g. PSI, and (b) SBAS InSAR stack.

SBAS was firstly proposed in (Berardino et al., 2002). Several similar techniques exist, including (Mora et al., 2003; Blanco-Sánchez et al., 2008), an update from the original group (Lanari et al., 2004), as well as (Goel and Adam, 2012a) which make use of adaptive multi-looking to further reduce the decorrelation of DSs — an improvement following the innovation of SqueeSAR. Nevertheless, the spirit of SBAS is to use small baseline, which does

not only limit to DS. This has been shown in applying to partially coherent PSs (Wegmüller et al., 2010; Perissin and Wang, 2012).

A generally recognized procedure of SBAS is as follows (Berardino et al., 2002).

- (1) Given a stack of  $N+1$  co-registered images, multiple interferograms of small spatial and temporal baselines with individual masters are formed, instead of  $N$  interferogram with a single master. The atmospheric phase is assumed to be already removed. Otherwise, the procedure is similar to that explained in PSI.
- (2) Coherent pixels are selected by thresholding on the coherence. Note its difference to PS selection. The coherence explicitly defines the variance of interferometric phase noise by the analytical expression (Tough et al., 1995; Bamler and Hartl, 1998):

$$\sigma_{\varphi}^2(|\gamma|) = \pi^2/3 - \pi \arcsin(|\gamma|) + \arcsin^2(|\gamma|) - \text{Li}_2(|\gamma|^2)/2 \quad (3.7)$$

where  $\text{Li}_2$  is the dilogarithm.

- (3) The interferograms are unwrapped in 2-D. The small baselines reduce the topographic phase contribution, and eases the unwrapping step.
- (4) The topography height and the low frequency deformation signal (usually a linear deformation model) are estimated. This step is similar to the least square estimation in LAMBDA after the  $2\pi$  ambiguity been fixed. The phase contribution from topography is subtracted from the unwrapped phase. Considering a possible phase unwrapping error in step 3, the residual phase is usually unwrapped once again.
- (5) The remaining interferometric phase has contributions from non-linear motion signal and noise. They are with respect to multiple master images. In order to reconstruct the  $N$  absolute phase with respect to a single master, one needs to solve a linear equation system  $\Delta\boldsymbol{\varphi} = \mathbf{A}\boldsymbol{\varphi} + \boldsymbol{\varepsilon}$ , i.e.:

$$\begin{bmatrix} \varphi_{2-1} \\ \varphi_{3-2} \\ \vdots \end{bmatrix} = \begin{bmatrix} -1 & 1 & \cdots \\ \vdots & -1 & 1 \\ & -1 & 1 \\ & & \ddots \\ & & & -1 & 1 \end{bmatrix} \begin{bmatrix} \varphi_1 \\ \varphi_1 \\ \vdots \\ \varphi_N \end{bmatrix} + \boldsymbol{\varepsilon} \quad (3.8)$$

where  $\Delta\boldsymbol{\varphi} \in \mathbb{R}^M$  is the unwrapped phase of the small baseline interferograms after removing the topographic phase,  $\mathbf{A}$  of dimension  $M \times N$  is the design matrix of the small baseline interferograms,  $\boldsymbol{\varphi} \in \mathbb{R}^N$  the absolute phase with respect to a single master, and  $\boldsymbol{\varepsilon}$  the phase noise.

Equation (3.8) is often overdetermined but ill-posed, due to the fact that not all the interferogram are connected to a single subset, rendering the rank of  $\mathbf{A}$  to be less than  $N$ . Therefore, regularization is required. Singular value decomposition (SVD) has been used for this task (Liebhart et al., 2010).

## Summary

The parameter estimation in SBAS is all based on linear least squares. Hence similar to LAMBDA estimator, it also assumes a Gaussian phase noise, which is not true under the strict DS model. Regularization is required if the interferograms do not connect into a single subset. Computational-wise, SBAS requires the unwrapping of a substantial number of interferograms, which is usually much more than  $N$ . Therefore, the phase unwrapping is the most time consuming and risky step in SBAS.

### 3.1.3 SqueeSAR (Adaptive Multilooking)

#### General Remark

Same as SBAS, SqueeSAR also exploits the DSs. Instead of restricting the DS decorrelation using interferograms of small baselines, SqueeSAR tackles it by averaging the DS pixels. As we know, the MLE of the interferometric phase given  $M$  ergodic DS samples is

$$\hat{\phi}_{MLE} = \arg \left( \sum_{m=1}^M v[m] \right) \quad (3.9)$$

This idea is also known as *multilooking*, e.g. the "box car" filter. However, the samples selected in a rectangle window are often not ergodicity, and hence, produce unsatisfactory results. Therefore, the core innovation of SqueeSAR (Ferretti et al., 2011) and its variants (Wang et al., 2012; Goel and Adam, 2012a; Jiang et al., 2015) is adaptive multilooking.

A generally recognized SqueeSAR workflow is as follows (Ferretti et al., 2011):

- (1)  $N$  interferograms are formed with respect to a single master. This step is the same as PSI.
- (2) The neighbourhood (similar pixels) of each pixel is found by comparing their intensity series using statistical test. The neighbourhood pixels are averaged to produce the filtered interferogram. The statistical test can be either parametric or non-parametric.
  - ◇ Non-parametric test

#### *Kolmogorov-Smirnov (KS) test*

It measures the maximum distance between the empirical cumulative distribution functions (CDFs) of two data sets (Ferretti et al., 2011), i.e.

$$D_{KS} = \max_x |F_a(x) - F_b(x)| \quad (3.10)$$

where  $F_a(x)$  and  $F_b(x)$  are the empirical CDFs of the two target pixels' intensities, respectively. By setting a threshold on the distance, the null hypothesis that the two pixels belong to the same distribution is accepted or rejected. However, the maximum distance of two CDFs usually happens in the midway of the curve for any distribution. Therefore, the KS test is not sensitive to unknown distributions with variations on the head or tail.

#### *Anderson-Darling (AD) test*

Likewise, AD test also measures the distance between two CDFs, but it includes weighting according to the shape of the distribution (Parizzi and Brcic, 2011; Wang et al., 2012; Goel and Adam, 2012a), i.e.

$$A^2 = \frac{N}{2} \sum \frac{(F_a(x) - F_b(x))^2}{F_{a,b}(x)(1 - F_{a,b}(x))} \quad (3.11)$$

where  $F_{a,b}(x)$  is the CDF of the combined pixels' intensities. It places more weight on the head and tails of the distributions. AD test is actually a weighted  $L_2$ -norm of the difference of two CDFs.

- ◇ Parametric test

Parametric test compares the intensity series of two pixels through generalized likelihood ratio test conditioned on the null and alternative hypotheses, with a given

parametric model of the intensity distribution. It gives the best performance if the data follows the assumed model.

The marginal distribution of a DS intensity series, if not considering the correlation between images, is  $\chi^2$  distribution with two DoF (equation (2.3)). Since the expected intensity fully characterizes this distribution, the null and alternative hypotheses are  $H_0: \bar{I} = \bar{I}_a = \bar{I}_b$  and  $H_1: \bar{I}_a \neq \bar{I}_b$  where  $\bar{I}_a$ ,  $\bar{I}_b$ , and  $\bar{I}$  are the expected intensity of target pixel  $a$ ,  $b$ , and combined. The test statistic is then (Parizzi and Brcic, 2011):

$$T = 2N \ln(\hat{\bar{I}}) - N \ln(\hat{\bar{I}}_a) - N \ln(\hat{\bar{I}}_b) \stackrel{H_1}{>} 0 \quad (3.12)$$

where  $\hat{\bar{I}}_a$ ,  $\hat{\bar{I}}_b$ , and  $\hat{\bar{I}}$  are the MLE of  $\bar{I}_a$ ,  $\bar{I}_b$ , and  $\bar{I}$ , respectively.

An advance of the parametric similarity test is demonstrated in the non-local InSAR framework (Deledalle et al., 2015), where the similarity is compared patch-wisely taken into account the interferometric phase.

- (3) The multilooked interferometric phase is not conservative, i.e.  $\bar{\varphi}_{1-2} + \bar{\varphi}_{2-3} + \bar{\varphi}_{3-1} \neq 0$ . They are "triangulated" by solving a linear equation system similar to the SBAS equation (3.8).
- (4) The phase history parameters of filtered DS are estimated as a PS using periodogram or LAMBDA.

### DS MLE

On the estimation point of view, the MLE of the phase of CCG distributed DS can be formulated as follows (De Zan and Rocca, 2005; De Zan, 2008; Guarnieri and Tebaldini, 2008; Ferretti et al., 2011; Wang et al., 2012):

$$\hat{\theta} = \arg \min_{\theta} \left\{ \mathbf{g}^H \mathbf{\Phi}(\theta) \text{abs}(\hat{\mathbf{C}})^{-1} \mathbf{\Phi}(\theta)^H \mathbf{g} \right\} \quad (3.13)$$

where  $\mathbf{\Phi}(\theta) \in \mathbb{C}^{N \times N}$  is the diagonal matrix containing the modelled phase of  $\mathbf{g}$ , and  $\text{abs}(\cdot)$  is the element-wise absolute value of a matrix. Given a stationary neighbourhood of the target DS, the MLE maximize the joint likelihood:

$$\hat{\theta} = \arg \min_{\theta} \sum_{m=1}^M \mathbf{g}_m^H \mathbf{\Phi}(\theta) \text{abs}(\hat{\mathbf{C}})^{-1} \mathbf{\Phi}(\theta)^H \mathbf{g}_m \quad (3.14)$$

where  $m$  is the index of the neighbouring sample. In (3.14), the same phase parameters  $\theta$  is assumed for all the observations, i.e.  $\mathbf{g}_i$  for  $m = 1, \dots, M$ .

### Summary

SqueeSAR is a powerful method to make use of partially decorrelated DSs. The core of SqueeSAR is adaptive averaging. SqueeSAR represents the state-of-the-art in DS processing, since it takes both advantage of interferograms of small baselines, as well as the adaptive multilooking.

The performance of standard SqueeSAR greatly depends on the similarity test. The performance of similarity tests degrades as the number of images decreases (Parizzi and Brcic, 2011). Computational-wise, SqueeSAR is tens of times more expensive than PSI. However, the overall computation is still moderate compared to TomoSAR.

## 3.1.4 SAR Tomography and Differential SAR Tomography

### General Remark

As mentioned in section 2.1.3, TomoSAR solves the layover problem where the full reflectivity profile  $\gamma(s)$  along the third dimension *elevation*  $s$  is retrieved. The following equation is obtained by subtracting the phase at zero elevation from equation (2.4) (ignore the second order elevation term in the phase) (Reigber and Moreira, 2000; Lombardini et al., 2003; Fornaro et al., 2003; Zhu, 2008):

$$g_n = \int_{\Delta s} \gamma(s) \exp\left(-j \frac{4\pi b_n}{\lambda R} s\right) ds \quad (3.15)$$

where  $g_n$  and  $b_n$  are the pixel value and the perpendicular baseline of the  $n$ th image. In the case of differential TomoSAR (D-TomoSAR), equation (3.15) is extended into higher dimensions (Zhu and Bamler, 2010b, 2011). Write (3.15) in a more compact matrix form, it is

$$\mathbf{g} = \mathbf{R}\boldsymbol{\gamma} \quad (3.16)$$

where  $\mathbf{R}$  is basically a discrete Fourier matrix whose frequencies are linearly proportional to the baselines. A generally recognized TomoSAR workflow is as follows (Zhu and Bamler, 2010b):

- (1) Interferograms are formed like PSI. The atmospheric phase screen is mitigated.
- (2) The reflectivity profile is reconstructed using spectral estimation techniques.
- (3) The scatterers' positions and motion parameters are determined by detecting maximum on the reflectivity profile.

Since the first step is the same as standard PS processing, the following discussion will focus on the estimation of the reflectivity profile.

#### MLE and Maximum A Posteriori (MAP) Estimator

Given a deterministic  $\mathbf{g}$  with i.i.d. complex Gaussian noise, the MLE of  $\boldsymbol{\gamma}$  is

$$\hat{\boldsymbol{\gamma}}_{MLE} = \min_{\boldsymbol{\gamma}} \|\mathbf{g} - \mathbf{R}\boldsymbol{\gamma}\|_2^2 \quad (3.17)$$

For stochastic  $\mathbf{g}$ , the MLE should consider the covariance matrix of the signal (Wax, 1991; Sauer et al., 2009; Tebaldini, 2010). Since the dimension of  $\boldsymbol{\gamma}$  is often much larger than  $\mathbf{g}$ , the equation  $\mathbf{g} = \mathbf{R}\boldsymbol{\gamma}$  is an underdetermined system. There are a few methods to regularize the equation system, i.e. MAP estimators:

- ◇ Nonlinear least square (Stoica and Moses, 2005): Assuming the resolution cell consists  $K$  scatterers with their elevation  $\mathbf{s} = [s_1, s_2, \dots, s_K]$ , this system can be reduced to where the support of the signals are. For small number of  $K$  which is usually the case, it becomes an overdetermined system.

$$\hat{\boldsymbol{\gamma}}(\hat{\mathbf{s}})_{NLS} = \min_{\boldsymbol{\gamma}(\mathbf{s})} \|\mathbf{g} - \mathbf{R}(\mathbf{s})\boldsymbol{\gamma}(\mathbf{s})\|_2^2 \quad (3.18)$$

The dimension of  $\mathbf{R}(\mathbf{s})$  and  $\boldsymbol{\gamma}(\mathbf{s})$  are much smaller than  $\mathbf{R}$  and  $\boldsymbol{\gamma}$ , respectively. However, they depend on the location of  $\mathbf{s}$  which needs to be estimated simultaneously. Application in layover separation has been demonstrated in (Ferretti et al., 2005; Zhu, 2008).

- ◇ Tikhonov regularization: apart from the undermined equation system, the matrix  $\mathbf{R}$  can be rank deficient due to the irregularity and small span of the baseline. That is to say, (3.17) can be ill-posed and underdetermined. The Tikhonov regularized MAP is as follows:

$$\hat{\boldsymbol{\gamma}}_{Tikho} = \min_{\boldsymbol{\gamma}} \|\mathbf{g} - \mathbf{R}\boldsymbol{\gamma}\|_2^2 + \alpha_{Tik} \|\boldsymbol{\gamma}\|_2^2 \quad (3.19)$$

which restricts the energy of  $\boldsymbol{\gamma}$  by performing  $L_2$ -norm regularization. (Zhu and Bamler, 2010b) implemented the MAP using SVD with Wiener weighing on the eigenvalues. Alternatively, (Fornaro et al., 2003) truncated small eigenvalues.

- ◇ Compressive sensing (CS)-based estimators exploits the sparsity prior of the reflectivity profile, i.e. assuming only small number of scatterers on the reflectivity profile. Instead of the  $L_2$ -norm regularization in equation (3.19), sparse reconstruction of  $\gamma$  penalize on its  $L_1$ -norm:

$$\hat{\gamma}_{CS} = \min_{\gamma} \|\mathbf{g} - \mathbf{R}\gamma\|_2^2 + \alpha_{CS}\|\gamma\|_1 \quad (3.20)$$

The  $L_2$ - plus  $L_1$ -norm minimization is a sparse reconstruction problem. Its first application in TomoSAR was presented in (Zhu and Bamler, 2009) using real data. The same authors also developed the promising SL1MMER algorithm, which delivered the first CS-based TomoSAR reconstruction of single building (Zhu and Bamler, 2010a). The SL1MMER algorithm was also recently extended to the M-SL1MMER (Zhu et al., 2015b) which exploits the group sparsity in the urban environment. M-SL1MMER achieves comparable result with far less images than SL1MMER and other existing algorithms. Other CS-based TomoSAR with COSMO-SkyMed data (Budillon et al., 2011) and airborne millimetre-wave SAR data (Schmitt and Stilla, 2013) were also presented. The CS-based TomoSAR algorithms are much more computationally expensive than the Tikhonov regularization.

### Summary

TomoSAR is so far the most competent InSAR method for urban and forest area monitoring. The high computational cost limits it for an extensive use like PSI, especially for the CS-based TomoSAR algorithms.

Since the forward model  $\mathbf{R}$  is a discrete Fourier transform, TomoSAR is basically a spectral estimation problem. Many classical algorithms can be applied to TomoSAR, such as beamforming (Pasquali et al., 1995; Lombardini et al., 2003; Fornaro et al., 2009), CAPON (Lombardini et al., 2001), MUSIC (Gini and Lombardini, 2002; Guillaso and Reigber, 2005; Huang et al., 2012; Nannini et al., 2012), etc. Please refer to (Stoica and Moses, 2005; Zhu, 2011) for an extensive review of these classical methods.

## 3.1.5 InSAR Point Cloud Fusion

### General Remark

The abovementioned InSAR techniques deliver 3-D point clouds together with the motion parameters. Since the point clouds are relative to their reference point, they need to be co-registered when considering the results from multiple stacks. Although the general point cloud fusion is a classical topic in the computer vision field, there is actually only a handful of studies on InSAR point cloud fusion, especially for point clouds from image stacks of cross-heading orbits (i.e. ascending and descending), because there is theoretically no corresponding point from such two point clouds.

The first attempt of fusing cross-heading InSAR point clouds in urban area was presented in (Gernhardt and Bamler, 2012). This algorithm is essentially a 3-D surfaces matching. After removing the façade points where the point correspondence is most unlikely to appear, the point correspondences are found by searching closely spaced point pairs on surface. Therefore, this algorithm does not address the exact point correspondence.

The fusion of along-heading (both ascending or both descending) point clouds or point clouds of smooth terrain is less challenging. Classical point cloud co-registration methods such as iterative closest point (ICP) can be directly applied. (Gernhardt et al., 2012) has demonstrated the direct application of ICP on multiple InSAR point clouds of a volcano.

### Summary

The major difficulty in cross-heading InSAR point clouds fusion is the lack of exact corresponding point pairs, due to the SAR side-looking imaging geometry. The current methods

are all based on inexact match, except a recent advance (Zhu et al., 2014, 2015c) where the authors manually locate dozens of point pairs of street lampposts which are supposed to be the closest match to exact point pair, and performed careful calculation of the absolute position of the lamppost using the concept of StereoSAR (Gisinger et al., 2015) taking into account the diameter the of the lamppost.

### 3.1.6 Optical and SAR Images Fusion

#### General Remark

There have been many works regarding the fusion of optical and SAR images. As (Soergel, 2010) mentioned: "SAR inevitably requires an oblique scene illumination resulting in undesired occlusion and layover especially in urban areas", optical images can complement such disadvantage of SAR images. The fusion follows basically three categories: 1) geometric fusion, 2) joint height reconstruction, and 3) joint classification.

#### Geometric co-registration

Geometric fusion is the basis of most of the higher level fusion algorithms. The current techniques are mostly based on matching of 2-D spatial features, e.g. contours, edges, templates, etc. (Li et al., 1995; Ali et al., 2002; Cheng et al., 2004; Hong and Schowengerdt, 2005; Fan et al., 2013; Palubinskas and Reinartz, 2015). They are good for large scale analysis, but not applicable for the analysis of individual buildings, especially for high resolution data.

A precise 3-D model is crucial in co-registration of SAR and optical images, for example (Auer et al., 2015) used carefully reconstructed 3-D models from photogrammetric measurements; (Tao et al., 2014) used a precise airborne LiDAR point cloud; and (Schack et al., 2015) employed a PS point cloud retrieved from tens of high resolution TS-X spotlight images. Without a precise 3-D model, one needs to assume a strict prior (Brunner et al., 2010), such as rectangle footprint etc. Considering the complex 3-D geometry of urban area, there has not been an effective algorithm of matching SAR and optical image yet.

#### Building height reconstruction

Several literatures also explored the potential of 3-D reconstruction using a few SAR and optical images. The joint 3-D reconstruction can be said to be contextually or geometrically. Contextually, (Tupin and Roux, 2005) used the optical image as an a priori term in order to regularize the height reconstructed from radargrammetry. Geometrically, (Sportouche et al., 2011) projected the building footprint extracted from the optical image into the SAR image, and solved building height geometrically. Alternatively, (Wegner et al., 2014) retrieved the height of buildings from SAR and optical images independently, and fused them together using least square adjustment. Apart from these, there are also works on building reconstruction from multi-aspect SAR images (Xu and Jin, 2007; Thiele et al., 2007).

#### Joint classification

Many studies have been dedicated to the joint classification of SAR and optical images. Most of the applications are on urban mapping and classification (Gamba and Houshmand, 2002; Pacifici et al., 2008; Wegner et al., 2009; Amarsaikhan et al., 2010; Poulain et al., 2011), since the optical and SAR image provide complimentary information in urban area. As the focus of the thesis is not on the classification method, the relevant literatures are only briefly listed for the readers' interest.

#### Summary

Geometric fusion of optical and SAR image is an important step, because many higher level fusion is based on a very precise geometric fusion. So far, there has not been a precise fusion of meter-resolution SAR and optical image in urban area yet.

## 3.2 Contribution of this Thesis

### 3.2.1 Computationally Efficient TomoSAR Inversion

The much higher computational cost of TomoSAR is the major factor of preventing it from replacing PSI as the workhorse of InSAR monitoring. Therefore, the first contribution of this thesis is an efficient TomoSAR inversion approach. The proposed method combines PSI and TomoSAR by performing a pre-classification of single and double scatterers instead of the state-of-the-art approaches that always perform full model order selection meaning inevitable full TomoSAR inversion for all the pixels. The computational time is reduced by a factor of more than fifty.

### 3.2.2 TomoSAR Point Cloud Fusion

The current cross-heading InSAR point cloud fusion methods are based on inexact point match. They are essentially 3-D surfaces matching that rely on removing the façade points where the point correspondence is most unlikely to appear. Therefore, an InSAR point cloud fusion method was proposed making use of the rich façade information in TomoSAR point clouds in order to derive the theoretically exact point correspondence. The proposed method is much more computationally efficient than the state-of-the-art, while achieving consistent results.

### 3.2.3 InSAR Optical Fusion

InSAR estimates and optical images can complement each other. However, the fusion of the two in urban areas is still a topic under research, because a precise geometric fusion of optical and SAR images of urban areas is still a challenge. Therefore, an effective algorithm of geometrically and semantically fusing InSAR estimates and optical images are developed. The algorithm performs 3-D geometric co-registration, image classification, and eventually semantic level analysis in InSAR estimate.

### 3.2.4 Robust Estimators

#### **Robustness w.r.t. Large Phase Error**

A systematic solution to optimally weight the InSAR observation with large phase errors in the multipass InSAR algorithms was developed, which has not been addressed so far. Experiments showed that the proposed algorithm increases the efficiency of the estimator by a factor of 7 to 35 compared to state-of-the-art estimators, when 40% of the images are contaminated with large phase errors.

#### **Robustness w.r.t. Non-Gaussian Scatterers**

The covariance estimation of non-Gaussian and nonstationary InSAR samples has not been addressed so far. Therefore, a covariance estimator that is robust against non-Gaussian samples and invariant with respect to non-stationary phase was developed. The proposed robust covariance estimator requires no flattening of the interferometric phase, and performs well even on extremely heavily tailed distributions.



## 4 Summary of the Work

The objectives of the thesis are addressed in five full paper peer-reviewed journal articles (one still under review). This chapter gives a brief summary of the articles including the following main contributions.

- ◇ First demonstration of the integration of PSI and TomoSAR for operational processing, which delivered a TomoSAR reconstruction of an entire city using meter-resolution SAR data.
- ◇ Computationally efficient fusion of two cross-heading TomoSAR point clouds
- ◇ First demonstration of semantic fusion of optical image and TomoSAR of an urban area.
- ◇ First systematic proposal of robust estimators for multipass InSAR techniques, which includes a robust estimator that can handle large phase offsets in the observations, as well as a robust covariance estimator that is invariant to the non-stationary phase.

### 4.1 Computationally Efficient Tomographic SAR Inversion

Appendix A explains the detailed procedure of combining PSI and TomoSAR for computationally efficient tomographic SAR inversion. It delivers tomographic reconstruction of an entire city using meter-resolution spaceborne SAR data. The reduction of computational cost credits to the following three points:

- (1) Pre-classification of single and double scatterers using non-parametric methods without any full TomoSAR inversion.
- (2) Use PS periodogram to solve the detected single scatterers, and pass only the detected double scatterers to TomoSAR algorithms, e.g. SVD-Wiener and SLIMMER.
- (3) Use estimates from PSI as the prior for TomoSAR inversion.

The right subfigure of Figure 4.1 shows the flowchart of the method in Appendix A. The most important is the layer right after the PSI processing, where pixels are pre-classified to no- ( $H_0$ ), single- ( $H_1$ ), or double-scatterer ( $H_2$ ). The left figure shows the algorithm of pre-classification of scatterers. Firstly, the pixels without scatterer are separated using penalized likelihood ratio test (Wang et al., 2014):

$$\frac{f(\mathbf{g}|H_0)}{f(\mathbf{g}|H_1)\exp(-AIC)} \frac{P_0}{P_{1,2}} \frac{H_0}{H_1} > 1 \quad (4.1)$$

where  $H_0$  and  $H_1$  likelihood of each pixel are compared, taking into account the Akaike information criterion (AIC) penalty (Zhu and Bamler, 2010b), and  $P_0$ ,  $P_{1,2}$  are the prior probabilities of pixels with no scatterer and with single or double scatterer(s), respectively. At this stage,  $H_1$  likelihood is used for pixels with single and double scatterers. But the detection of single and double scatterer(s) is almost 100% correct with 0% false alarm rate, because the  $H_1$  likelihood of the pixels with double scatterers are still much higher than their  $H_0$  likelihood.

#### 4.1.1 Pre-classification of Single and Double Scatterers

The pre-classification of pixels with single or double scatterer(s) is carried out using a few easily accessible quantities associated with each pixel. They are amplitude dispersion index (ADI), the likelihood to single scatterer model ( $H_1$ ), and the pixel mean intensity with the following three reasons:

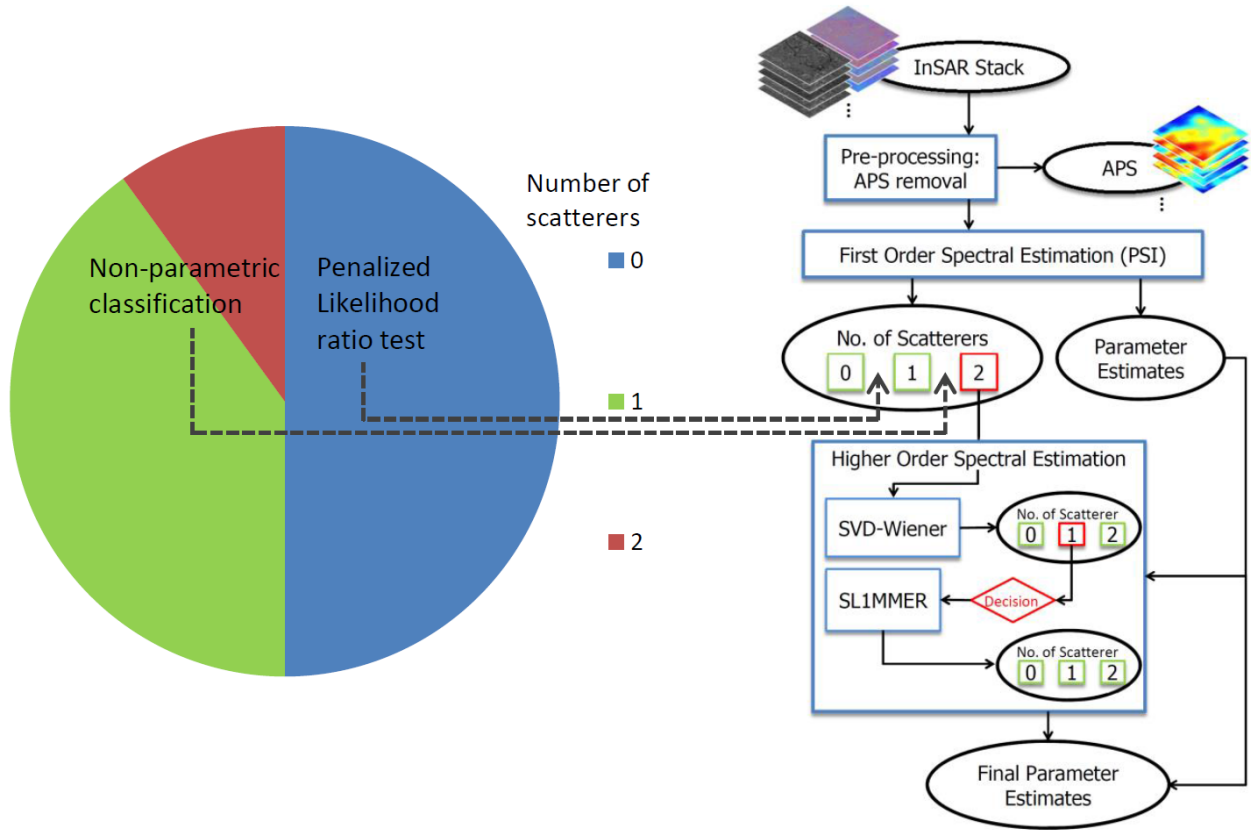


Fig. 4.1. left: Algorithm of pre-classification of non-, single-, and double-scatterers. Non-scatterers are detected by penalized likelihood ratio test. Single and double scatterers are classified using fast non-parametric methods without TomoSAR inversion, e.g. classification using mean intensity, amplitude dispersion index, etc. And right: the flowchart of the efficient TomoSAR inversion algorithm. The most important is the layer right after the PSI processing, where pixels are pre-classified, and PSI estimates are passed to TomoSAR processing as prior knowledge.

- (1) pixels with single scatterer experience lower ADI, since the amplitudes of pixels with only a single dominant scatterer do not show pronounced fluctuations as a function of baseline;
- (2) the  $H_1$  likelihood tends to be lower if multiple scatterers are present in one resolution cell;
- (3) the abovementioned two aspects are based on a reasonable SNR range. Therefore, only relatively bright pixels should be selected as candidates of pixels with double scatterers.

As an example, Figure 4.2 shows the joint distribution of mean intensity and the  $H_1$  log-likelihood of single and double scatterers, respectively. Here, we must note that the real ground truth of single and double scatterers classification is not obtainable. The single and double scatterers in the following plots refer to those detectable ones from the estimation point of view. In Figure 4.2, they are detected by the CS-based SL1MMER algorithm, which is one of the state-of-the-art algorithms. Based on these, it can be observed that the  $H_1$  log-likelihood (represented by normalized residual sum of squares (RSS)) of the single scatterers shows more linear relationship with the pixel intensity, whereas they are more independent for double scatterers. Double scatterers are in general 20dB brighter than single scatterers. Different behaviour can also be observed in the joint distributions of the other quantities among the three.

Figure 4.3(a) shows the pre-classification result of single and double scatterers using the three quantities, with red representing pixels with double scatterers, green for single scatterers, and blue for no scatterer. A linear support vector machine (SVM) was employed as the classifier. For comparison Figure 4.3(b) shows the classification result from the CS-based method SL1MMER (Zhu and Bamler, 2010a, 2012) as a reference, which is the best single

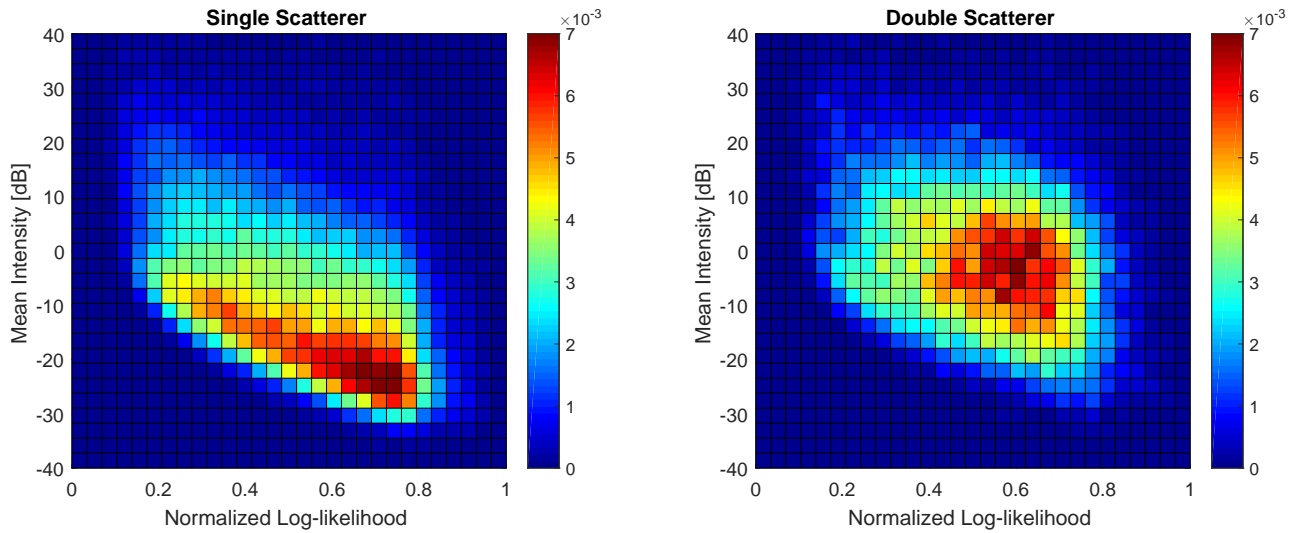


Fig. 4.2. The joint distribution densities of mean intensity and the log-likelihood to the single scatterer model for pixels with left: single scatterer, and right: double scatterer. The log-likelihood is represented by the residual sum of squares normalized to  $[0 \ 1]$ .

and double scatterer discrimination one can achieve so far. Figure 4.3(c) is the final classification after the higher order spectral estimation.

In Figure 4.3(a), 47% of the pixels are discarded as no-scatterers, 36% are detected as single scatterers, and the rest 17% are detected as double scatterers. Taking the SL1MMER result as the reference, the classification of pixel with no scatterer is almost 100% correct, and the detection rate of pixels with double scatterers is 72.4%, with a false alarm rate of 25.7% which is why much more pixels are detected as double scatterers. However, higher detection rate (consequently higher false alarm rate) is still favoured in the processing, because the falsely detected double scatterers can still be corrected in the higher order spectral estimation, but not for the non-detected ones. Depending on the processing power, the threshold can be shifted towards higher detection rate, affording higher false alarm rate, and therefore more processing effort by more frequently applying the SVD-Wiener and the SL1MMER.

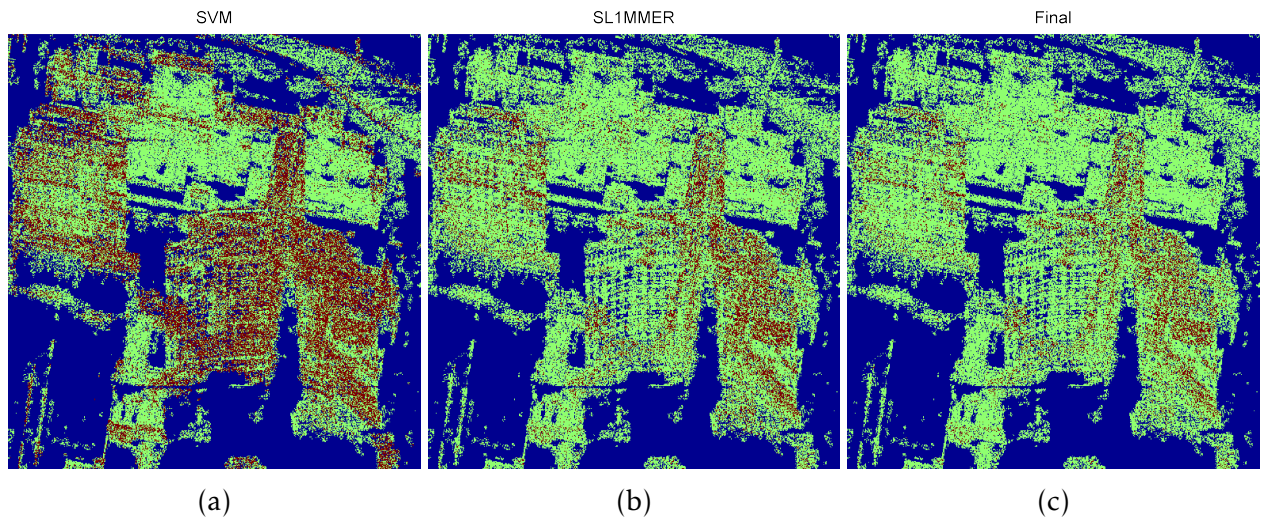


Fig. 4.3. (a) the result of the pre-classification method, (b) the result from the SL1MMER algorithm as a reference, and (c) the final classification after the higher order spectral estimation. The red pixels are detected double scatterers, the green are the single scatterers, and the blue are no scatterer.

Table 3. Overall computational cost of the proposed TomoSAR inversion method (using the test area shown in Figure 4.3 as the example), compared to using SL1MMER or SVD-Wiener only.

Processing Schemes	Percentage of pixels processed by each algorithm			
	Periodogram	SVD-Wiener	SL1MMER (full elevation search range)	SL1MMER (60m elevation search range)
100% SL1MMER (w/o prior)	—	—	100%	—
100% SVD-Wiener (w/o prior)	—	100%	—	—
Proposed method	100%	17%	—	9% (w/o prior)

#### 4.1.2 Computational Cost Analysis

Compare the PS periodogram and SVD-based TomoSAR MAP, the periodogram consists of multiplication of the measurement vector with the modelled phase, and repeats at different elevation positions. Therefore, the computational complexity is  $O(NL)$ , where  $N$  is the number of images (usually 20 - 50) and  $L$  is the discretization level in the elevation direction (usually 50 - 200 for TS-X). In the higher order spectral estimation, the SVD-Wiener needs at least  $O(N^2 + NL)$  according to the equation (11) in (Zhu and Bamler, 2010b). The CS-based TomoSAR algorithm requires an  $L_1$ - $L_2$  norm optimization which is extremely computationally costly, e.g. the well-known Basic Gradient Projection for Sparse Reconstruction (GPSR-Basic) solver (Figueiredo et al., 2007) requires at least  $O(KML_s^2)$ , where  $K$  is the number of iterations (approximately 20),  $M$  is the number of multiplication of a specific matrix in each iteration (modest, usually several), and  $L_s$  is the elevation discretization level required for sparse reconstruction, which is usually much higher than  $L$ . Our experience is the SL1MMER algorithm is several hundred times slower than SVD-Wiener.

When prior knowledge of the elevation span is available, the  $L$  can be restricted to a specific elevation range, and hence reduce the computational cost. For instance, assuming a uniform distribution of building heights prior over the processing area, the total processing cost can be reduced to half for SVD-Wiener when  $L \gg N$ , and to one third for SL1MMER. Therefore, it is always more economic to obtain the prior through some low-cost methods before the higher order spectral estimation.

The overall computational cost also depends on the percentage of pixels allocated to each algorithm. As an example, for the test area discussed, 100% of the pixels are first processed by periodogram, 17% are then passed to SVD-Wiener, and only 9% are finally processed by SL1MMER. This small percentage together with the restriction of the search space leads to an overall speed-up of about 50 times compared to SL1MMER-only processing. Table 3 compares the overall computational cost of the integrated inversion approach and the conventional TomoSAR-only inversion.

#### 4.1.3 Tomographic Reconstruction of Entire City

The abovementioned approach is applied on several stacks of TS-X high resolution spotlight images over Berlin. The following figure is the elevation estimates and a close-up view of one of the stacks which comprises 94 images. The proposed processing chain is able to catch most of the structure.



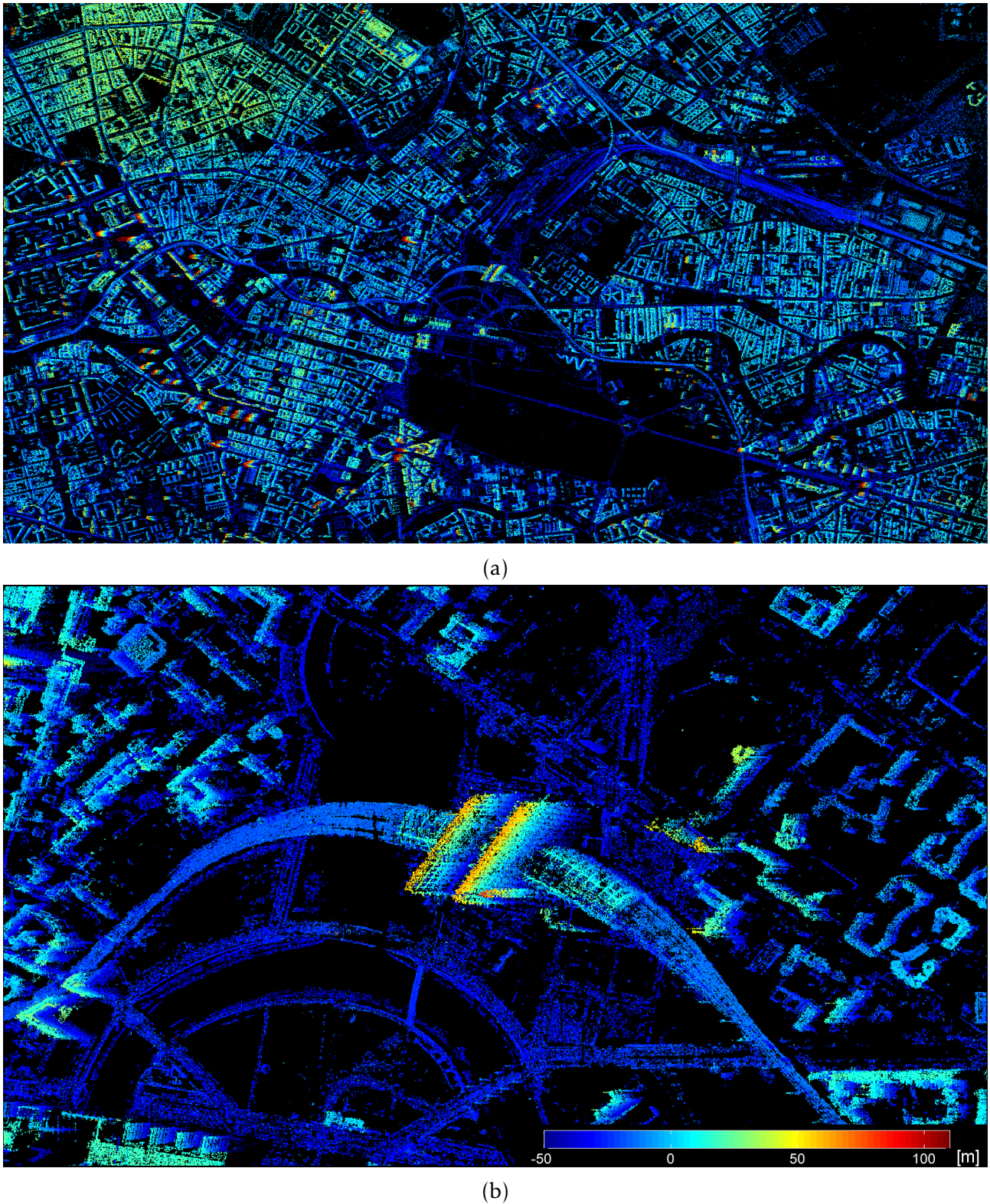


Fig. 4.4. (a) Elevation estimates of Berlin using the proposed approach, and (b) a close-up over the Berlin central station.

## 4.2 Automatic Feature-based TomoSAR Point Cloud Fusion

The previous section provides the solution to retrieve the 3-D TomoSAR point cloud of an entire city. A natural follow-up step is to combine several 3-D point clouds of the same area, especially for two cross-heading TomoSAR point clouds, i.e. derived from image stacks of ascending and descending orbits, as minimum of two TomoSAR point clouds from cross-

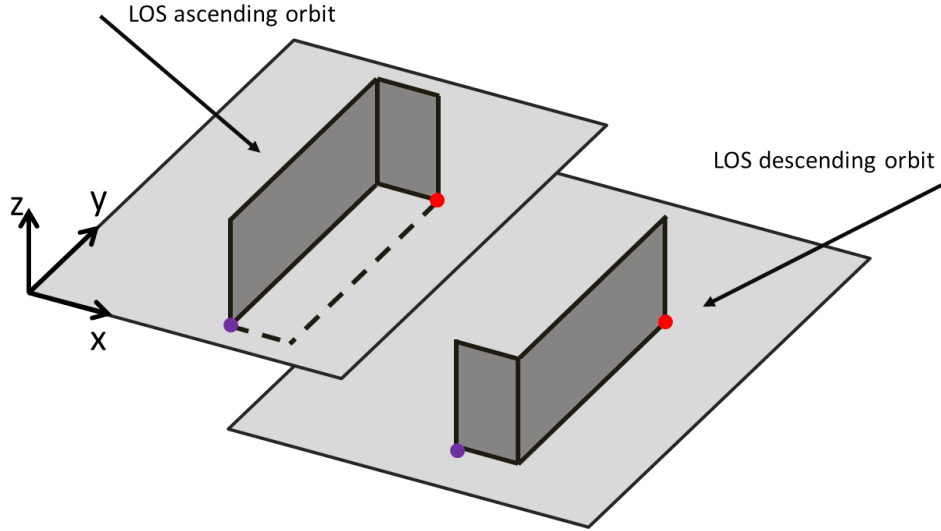


Fig. 4.5. Fusion geometry of the proposed L-shape method. The two building façades are coloured in dark grey representing the InSAR point clouds from cross-heading orbits whose line of sight directions are indicated by the two arrows. The light grey planes denote the corresponding ground level of the two point clouds. The two pairs of red and purple dots are the façade end point position to be estimated. Due to various facts, e.g. material, decorrelation, imaging geometry, etc., these end points may not be exactly from the scatterers located on the true end position of the façades, but they are still the closest match to the exact point correspondence in two cross-heading TomoSAR point clouds.

heading orbits are required to achieve a complete monitoring over the whole area due to the SAR imaging side-looking geometry.

InSAR point clouds are relative measurements whose reference points are usually unknown. The fusion task is to estimate the unknown 3-D shift. However, challenge arises at no exact point correspondence can be directly found among the natural scatterers in two cross-heading InSAR point clouds.

The contribution of Appendix B is finding and matching the point correspondences in two cross-heading TomoSAR point clouds. It is not difficult to realize that the end positions of façades where the two point clouds converge are the closest match to the exact point correspondences. The proposed point cloud matching algorithm is named as "L-shape detection and matching", because building façades often appear as L-shapes in InSAR point cloud. This concept is shown in Figure 4.5. The L-shape end points are indicated by the two pairs of red and purple dots. The two building façades coloured in dark grey represent the point clouds from ascending and descending orbits whose line of sight (LOS) directions are indicated by the two arrows, respectively. And the light grey planes denote the corresponding ground levels of the two point clouds.

Figure 4.6 shows the flowchart of the algorithm. Emphasis is put on the two most important procedures: point density estimation and L-shape detection which are marked as red in the figure.

#### 4.2.1 Point Density Estimation

Because of the SAR side-looking geometry, TomoSAR retrieves dense façade points. The estimation of point density is for detecting the façade points based on which the locations of the façade end points are estimated. Therefore, the point density estimation is the most crucial step in the whole algorithm.

We used a directional window for the density estimation, considering that façades are usually linear. This idea is illustrated in Figure 4.7. It shows the 2-D top view of a point cloud, with the color indicating the point density, and the red dot being the target point whose point density is to be estimated. The estimation follows three steps: firstly, points within

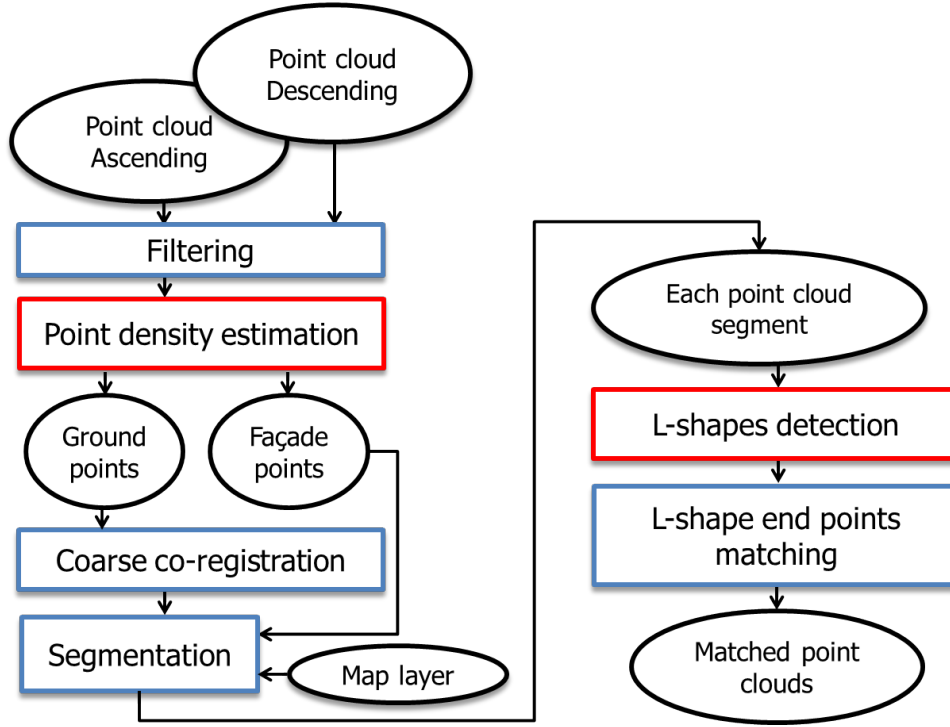


Fig. 4.6. Flowchart of the proposed algorithm. The black ellipses represent input/output data. The blue rectangles are procedures/processes.

a larger area (the red dashed rectangle) centered at the target point are selected; then a straight line segment (the green dashed line) passing through the target point is fitted using a robust algorithm, e.g. random sample consensus (RANSAC), robust principle component analysis (PCA), etc.; and lastly, only the inliers (within the solid red rectangle) of the line segment are counted for the point density, instead of counting all the points in the black rectangle.

The benefit of using adaptive directional window for point density estimation can be clearly seen in Figure 4.8 where subfigures (a) and (b) are the density estimates using directional and rectangular windows, respectively. It is obvious that the directional window emphasizes the façade points, while keeping the point density of non-façade points unchanged. And hence, it improves the detection rate of the façade points. The final façade points are selected by thresholding on the point density.

#### 4.2.2 L-shape End Point Estimation

Based on the detected façade points, a dominant L-shape is found for each building block by a weighted Hough transform (Wang and Zhu, 2015a). Since Hough transform only gives discretized estimates of the L-shape end points, their precise positions must be re-estimated.

The position of L-shape end point includes its horizontal 2-D coordinates and its height. The height is defined as the position where the façade intersects the ground plane, as depicted already in Figure 4.5.

The horizontal coordinates are estimated using a model-based approach. The procedures of the algorithm are briefly described as follows:

- (1) The two line segments of the L-shape are separated and rotated according to the façade direction. For example, one line of the L-shape with inlier points should look like Figure 4.9(a) where the black line is the estimated façade direction from the weighted Hough transform. Figure 4.9(b) is the rotated coordinates of the points in Figure 4.9(a). The middle section with much higher density belongs to the façade part.



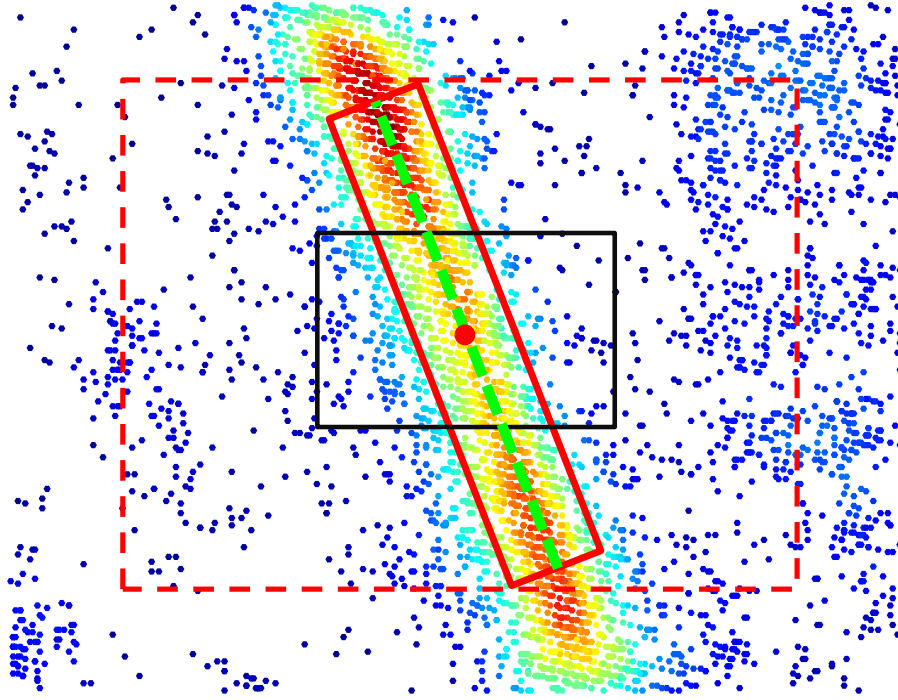


Fig. 4.7. Schematic drawing of point density estimation using directional window. The coloured points are the projection of the 3-D point cloud in horizontal plane, with the colour indicating the point density (blue: low density, red: high density). The green dashed line segment is a robust fit to the 2-D projection in the selected area within the red dashed rectangle. Points within the region bounded by the red solid rectangle are considered in the directional point density estimation, while those within the black rectangle are considered in the box-car density estimation.

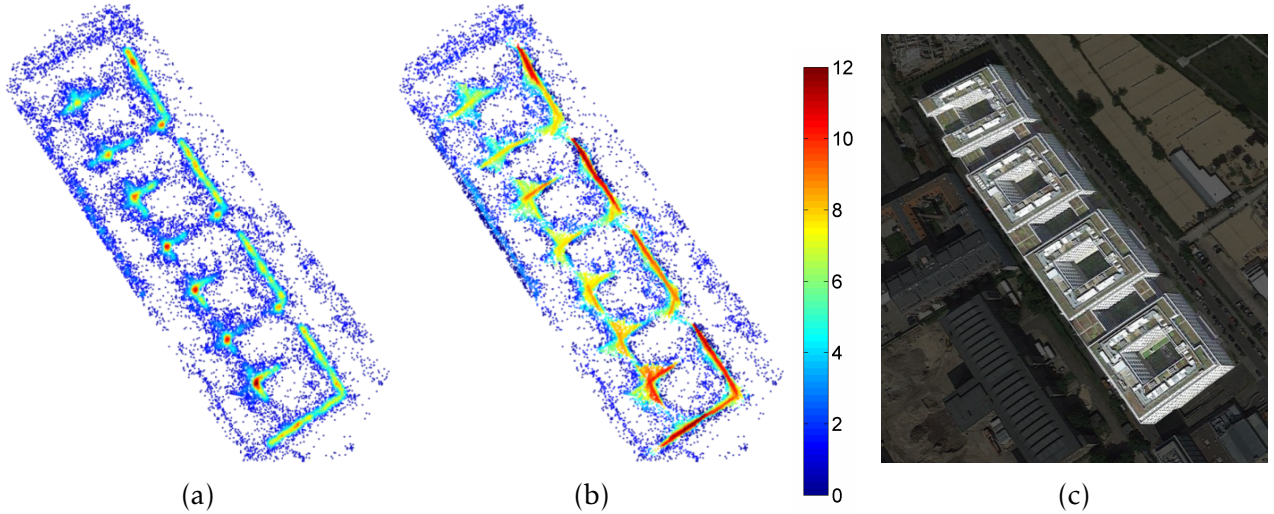


Fig. 4.8. Comparison of point density estimated using (a) directional window, and (b) rectangle window. The colour bar applies for both (a) and (b) with the unit  $[m^{-2}]$ . For reference, (c) is the optical image of the same area from Google. The size of directional and rectangle windows, in terms of area, are identical in this comparison. The directional window emphasizes the façade point, meanwhile keeps the point density of non-façade points unchanged compared to rectangle window. It improves the detection rate of façade points.

- (2) An estimate of the façade point density is obtained by convolving it with a rectangular window along the façade direction. For example, the blue curve in Figure 4.9(c) shows the point density estimate of the points shown in Figure 4.9(b).
- (3) The point density is modelled as a rectangle function plus a constant offset. Hence, the estimated point density is the convolution of two rectangle windows, which is a trapezoid. Therefore, the goal is to robustly deconvolve the trapezoidal façade density curve, which is equivalent to estimate the location of the two slopes of the trapezoid.
- (4) A moving line fitting method is developed to estimate the slopes' positions. A straight line is fitted to the points within a moving window with the same width as the rect-



angular window for density estimation. For any trapezoidal shape, high slopes should be detected at the position of the two slopes. By detecting the position of the two high slopes, one can determine the position of the two sides.

For robustness, the product of the absolute value of the slope and the number of inliers to the fitted line is considered in the detection of the two slopes, because the maximum number of inliers should be reached when the window centered at the left or the right slopes.

For example, Figure 4.9(d) shows the product of the absolute value of the slope and the number of inliers for the point density curve in Figure 4.9(c). Two prominent peaks can be detected. The positions of the peaks are the direct estimates of the center positions of the left and right position of the façade.

After obtaining the 2-D horizontal coordinates of the L-shape end point, we look for its neighbouring points on the ground level. The mean value of these points is an estimate of the height of the end point. A plane should be fitted to these points, if the local topography is not removed.

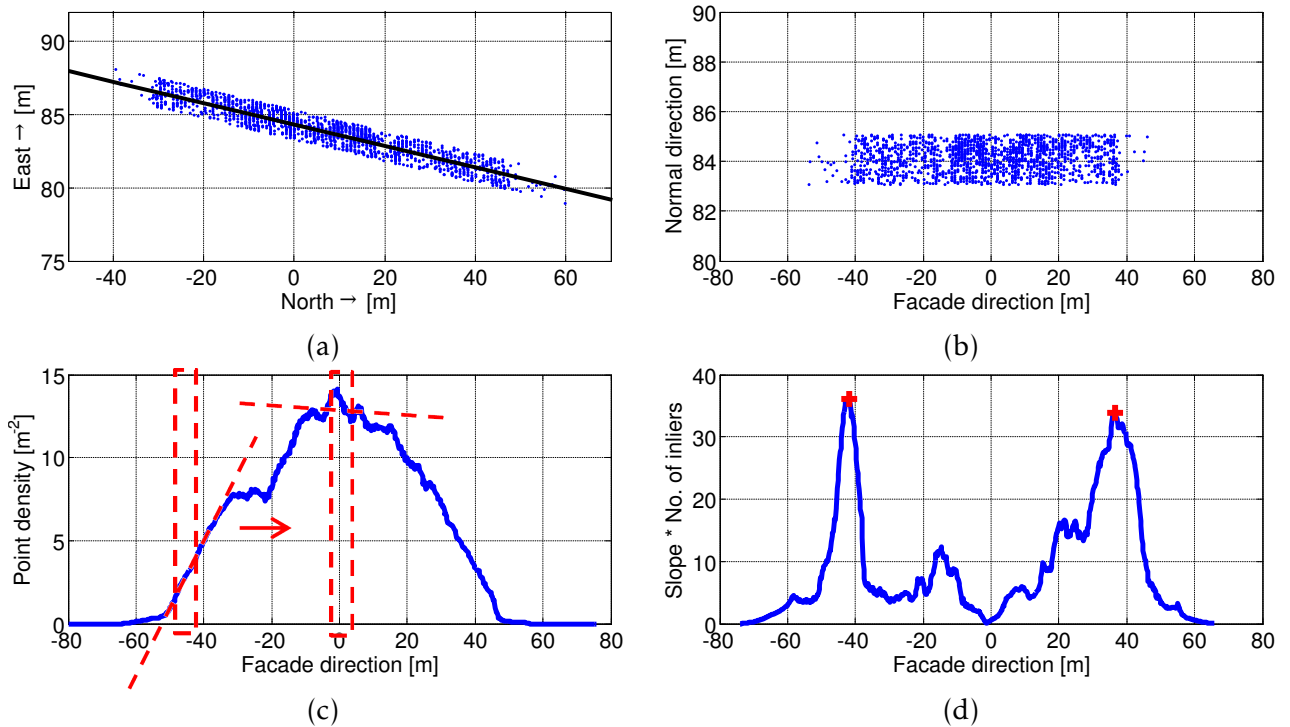


Fig. 4.9. (a) The inlier points of one line segment of an L-shape, with the black line being the façade direction obtained from the weighted Hough transform, (b) the coordinates of the inlier points projected to the façade direction, (c) the blue curve is the estimate of (b)'s point density using a rectangular window of 5m, and the red dashed rectangle and lines demonstrate the moving line fitting method, and (d) the product of the slope and the number of inliers of the fitted line using the points in the moving rectangle window along the point density estimate shown in (c). The two red crosses are the automatically detected façade start and end position at -41.38m and 37.22m, respectively.

### 4.2.3 Fusion Result

The proposed method was applied on two TomoSAR point clouds of Berlin contains about 20 million points each. Around 500 L-shapes are detected from each point cloud, corresponding to 1000 end points pairs. Figures 4.10(a) and (b) show the detected L-shapes and end points of the two point clouds, overlaying on their grey scale point density images, respectively. The matched L-shapes end points are shown in Figure 4.10(c). Closed quadrilateral implies both pairs of end points in an L-shape are matched. A single L-shape implies only one pair of end points in the L-shape is matched. In total, 150 pairs of end points are matched, which is sufficient to robustly estimate the 3-D shift.

This result is compared to the one computed using the method explained in (Gernhardt and Bamler, 2012). A difference of 0.51m in east direction; 0.09m in north direction and almost no difference in height is found. A 3-D view of the first fused TomoSAR point clouds of an entire city is shown in Figure 4.11. The height of the points is color-coded.

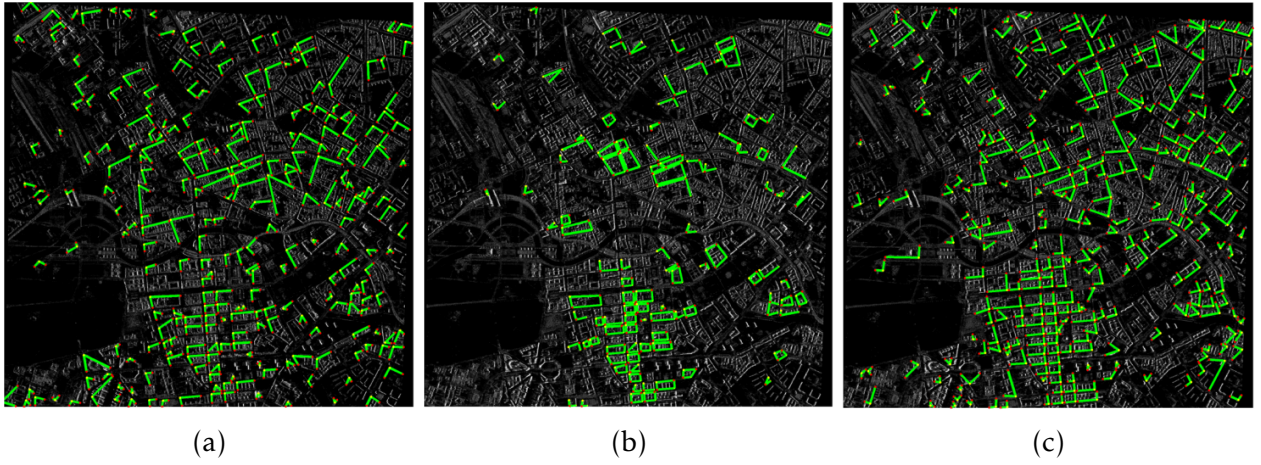


Fig. 4.10. (a) and (b): The detected L-shapes and their end points of the two input point clouds over Berlin, and (c) the matched L-shapes and end points. The L-shapes are plotted in green, and the end points are marked as red. They are overlaid on the grey scale point density image.

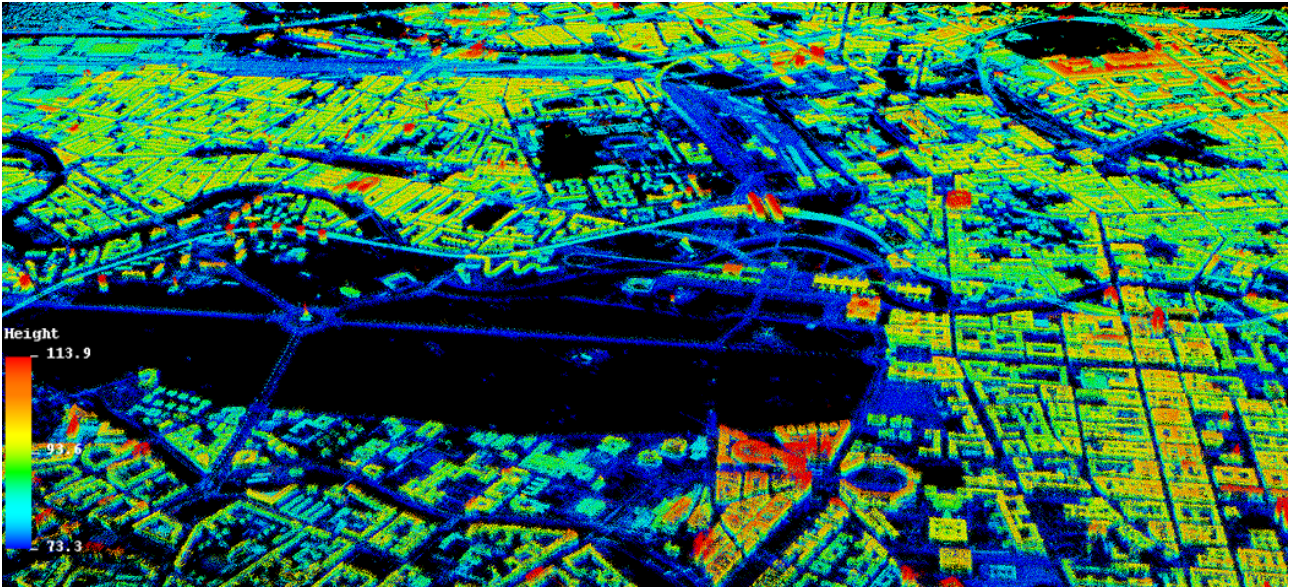


Fig. 4.11. The first fused TomoSAR point cloud of an entire city. The result combines an ascending and a descending point cloud of Berlin. The color indicates height on WGS84 reference surface. The unit of the colorbar is meter.

### 4.3 Semantic Interpretation of InSAR Estimates Using Optical Images

Having reconstructed the 3-D InSAR point cloud of an urban area, the scientific question of the origin of the scatterers in the real world arises. By fusing InSAR point clouds and optical images, we can trace the scatterers in the optical images. The interpretability of optical images also complements the InSAR deformation monitoring, permitting a systematic monitoring of urban areas by the semantics of infrastructures, e.g. monitoring of bridges, roads and railways. The contribution of Appendix C is a method of semantic interpretation of InSAR point cloud using optical images.



The basic idea is by geometrically 3-D co-registering optical images to an InSAR point cloud, we can transfer the semantics derived from the optical image to the InSAR point cloud for further analysis. The flowchart of the algorithm is as follows.

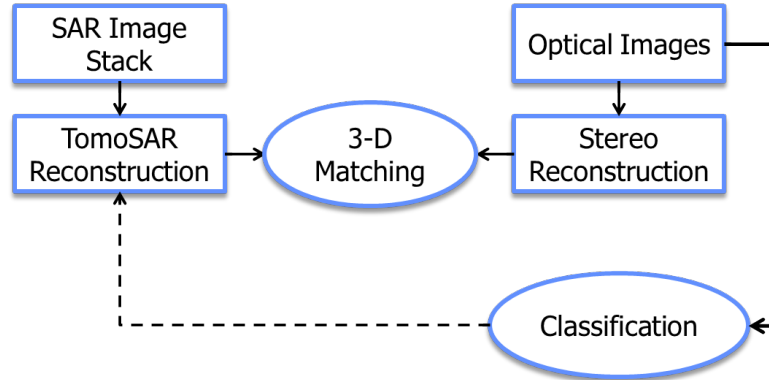


Fig. 4.12. Flowchart of the semantic InSAR point cloud interpretation method. The semantic classification of the InSAR point cloud is achieved by geometric co-registering the InSAR point cloud and the optical image to a reference model.

### 4.3.1 Geometric Co-registration of Optical Images and InSAR Point Cloud

The first contribution of the proposed algorithm is a strict 3-D geometric fusion of InSAR point cloud and optical images in urban area. SAR and optical images are different 2-D projections of a 3-D scene. Due to the complex urban topography, the co-registration must be done with a precise 3-D model. Figure 4.13 is a comparison of optical and SAR image of an urban area. A lot of details which are visible in the optical image are completely hidden underneath the strong reflection of the façade scatterers in the SAR image.

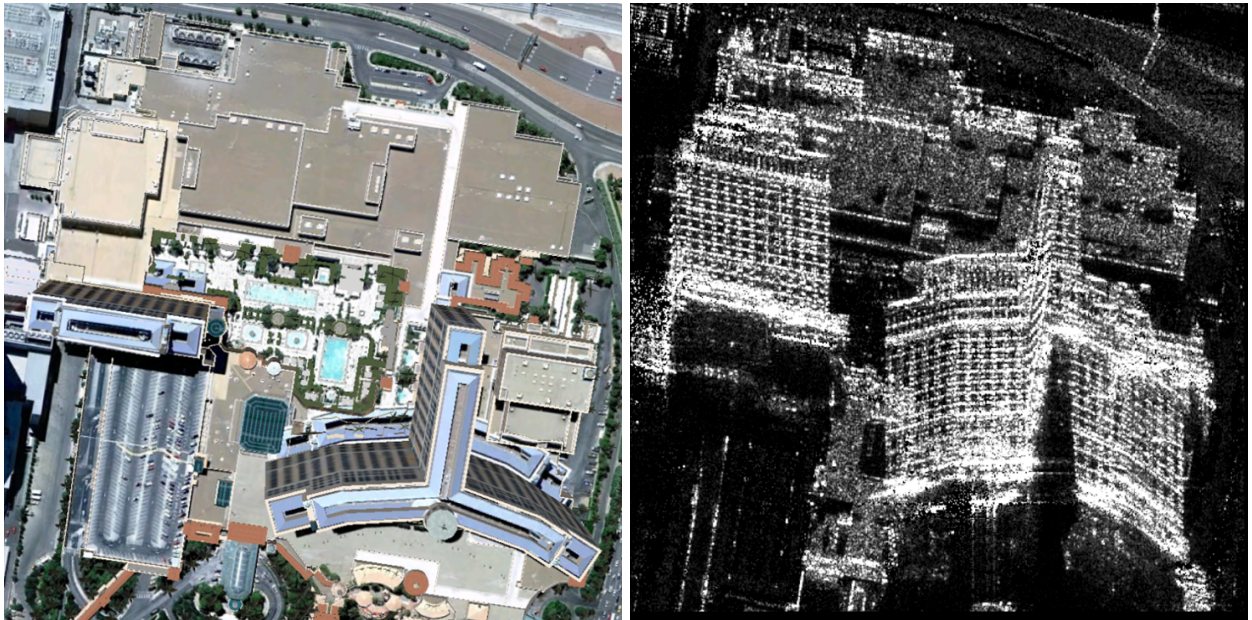


Fig. 4.13. Comparison of Google optical image (left) and TS-X high resolution spotlight image (right) of a same area. A lot of details visible in the optical image are completely hidden underneath the strong reflection of the façade scatterers in the SAR image. Images are from (Zhu and Bamler, 2012).

The proposed co-registration algorithm is as follows

- (1) Reconstruct 3-D point cloud from two or more optical images using stereo matching with structure from motion (SfM) if necessary
- (2) Reconstruct 3-D point cloud from SAR images using TomoSAR

- (3) 3-D co-registration of the optical and InSAR point clouds
- (4) Project optical image into the SAR geometry, a step also known as rendering

Since the first two steps are well explained in other literatures, the following text will focus on the co-registraion of optical and InSAR point cloud.

The aerial optical images are usually nadir-looking, in contrast to the side-looking geometry of SAR. In another word, façade points barely appear in optical point cloud while they are prominent in the TomoSAR point cloud. The point cloud derived from aerial optical images also in general has better relative positioning accuracy than the spaceborne TomoSAR point cloud. This difference is exemplified in Figure 4.14, where the left and right subfigures correspond to the TomoSAR and optical point clouds of the same area, respectively. These unique modalities have driven our algorithm to be developed in the following way:

- (1) Edge extraction
  - (a) The optical point cloud is rasterized into a 2-D height image.
  - (b) The point density of TomoSAR point cloud is estimated on the rasterized 2-D grid.
  - (c) The edges in the optical height image and the TomoSAR point density image are detected, respectively.
- (2) Initial alignment
  - (a) Horizontally by cross-correlating the two edge images.
  - (b) Vertically by cross-correlating the height histogram of the two point clouds.
- (3) Refined solution
  - (a) The façade points in both point clouds are removed.
  - (b) The final solution is obtained using *anisotropic* iterative closest point (ICP) with robustly estimated covariance matrix.

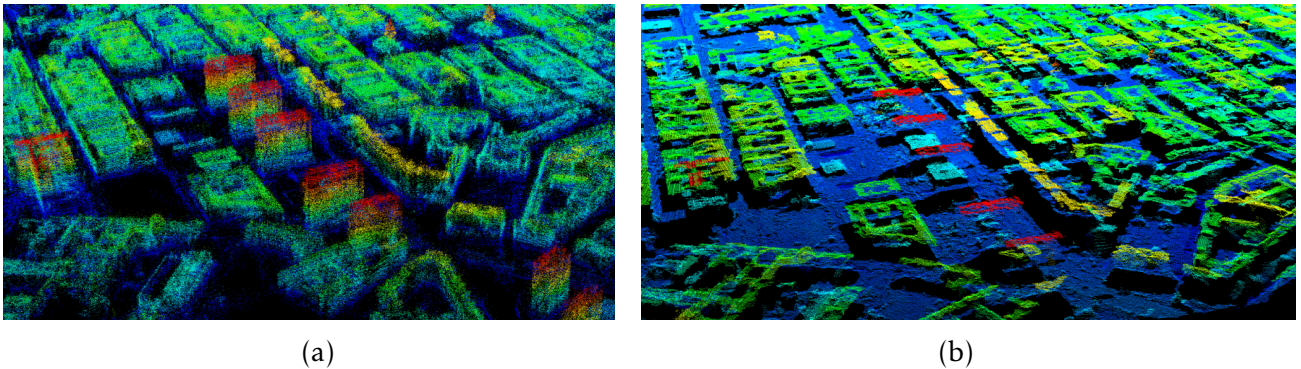


Fig. 4.14. (a) the TomoSAR point cloud of high-rise buildings, and (b) the optical point cloud of the same area. Building façades are almost invisible in the optical point cloud, while they are prominent in the TomoSAR point cloud.

More detail of the co-registration algorithm can be found in Appendix C. The final co-registration result is demonstrated in Figure 4.16. The top subfigure shows the reconstructed optical point cloud with colour representing the height, and the middle subfigure is the co-registered point cloud combining the optical one and two TomoSAR point clouds from ascending and descending viewing angles. Successful co-registration can be confirmed by the correct location of the façade points in the bottom of Figure 4.16 which is the top view of the fused point cloud with different colours representing different point clouds.

### 4.3.2 TomoSAR Point Cloud Rendering

Once the 3-D point clouds are co-registered, the relative positions of the optical image w.r.t. the SAR image are also solved. The optical image can be projected into the SAR geometry. This is also known as *rendering*: given a viewpoint, solve the occlusion problem. However, points do not occlude each other except degenerated points, because they are dimensionless.

Typically, the rendering of a point cloud requires reconstructing the 3-D surface and testing the depth of each point. But reconstructing reliable 3-D surface from TomoSAR point cloud is still a challenge task, although there are some pioneer works on reconstructing façades and roofs from spaceborne TomoSAR points (Zhu and Shahzad, 2014; Shahzad and Zhu, 2015). Therefore, the rendering technique should avoid the surface reconstruction.

This issue has been addressed in Appendix D by an improved hidden point removal operator (a point-based rendering technique). The hidden point removal operator (HPR) (Katz et al., 2007) inverts the original point cloud using certain function, so that the visible points will lie on the convex hull of the inverted point cloud. The most common inverting function is the spherical flipping. Given a point  $\mathbf{x}$  of any dimension, the *spherical flipping* function is defined as follows:

$$\hat{\mathbf{x}} = f(\mathbf{x}) = \mathbf{x} + 2(r - \|\mathbf{x}\|) \frac{\mathbf{x}}{\|\mathbf{x}\|} \quad (4.2)$$

where  $r$  is radius of the multi-dimensional sphere. The following figure illustrates the transformation function and HPR in 2-D. The blue dots are the original point cloud. And the red dots are the inverted one given a viewpoint  $O$ . The convex hull of the inverted point cloud is computed and the points lying on the convex hull are marked as visible.

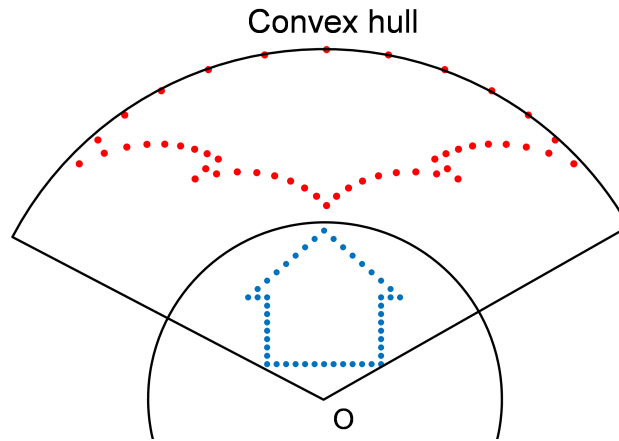


Fig. 4.15. Hidden point removal operator. The blue dots are the original point cloud. And the red dots are the inverted one given a viewpoint  $O$ .

Considering the much larger and isotropic noise in spaceborne TomoSAR point clouds than in typical computer graphics problems, the HPR is not directly applicable. Inspired by (Mehra et al., 2010), we introduced a visibility probability which is a Gaussian function of the point-to-convex hull distance  $d$ :

$$p(\hat{\mathbf{x}}) = \exp\left(-\frac{d^2}{2\sigma_{\hat{\mathbf{x}}}^2}\right) \quad (4.3)$$

where  $\sigma_{\hat{\mathbf{x}}}$  is the norm of the total noise standard deviations (all three directions) in the inverted point cloud.  $\sigma_{\hat{\mathbf{x}}}$  can be directly computed from the noise of the original point cloud.

Apart from this, we also include an image source consistency check by weighted majority voting to achieve consistent texturing among neighbouring points in the TomoSAR point cloud.

As a result, Figure 4.17(a) shows the first spaceborne meter-resolution 3-D TomoSAR point cloud of an urban area textured with colour from the optical images. Where no TomoSAR point is available, the black background is filled. The subfigure (b) is the same TomoSAR

point cloud, but colour-coded by its height. The dashed ellipses in subfigure (a) mark the Berlin central station and the main building of Universitätmedizin Berlin. The former consists of some railway segments on the ground level and the station building of about 5 stories tall, and the latter is a high-rise building of about 20 stories tall. The close up views of these areas are shown in subfigures (c) and (d). The boundary of the railway and the building edges are clearly projected. For the more challenging task of texturing high-rise objects, the proposed algorithm also successfully captures the regular window raster on the façade of Universitätmedizin Berlin. This implies a good estimation of the cameras parameters, of the relative positions between the cameras and the SAR sensor, as well as a good estimation of the points visibilities to the different cameras.

Of course, each point in the point cloud also contains its deformation parameters. By associating the deformation parameters with its optical appearance, we have an intuitive way of analysing the scattering mechanism of different objects. This will be exemplified in the next section.



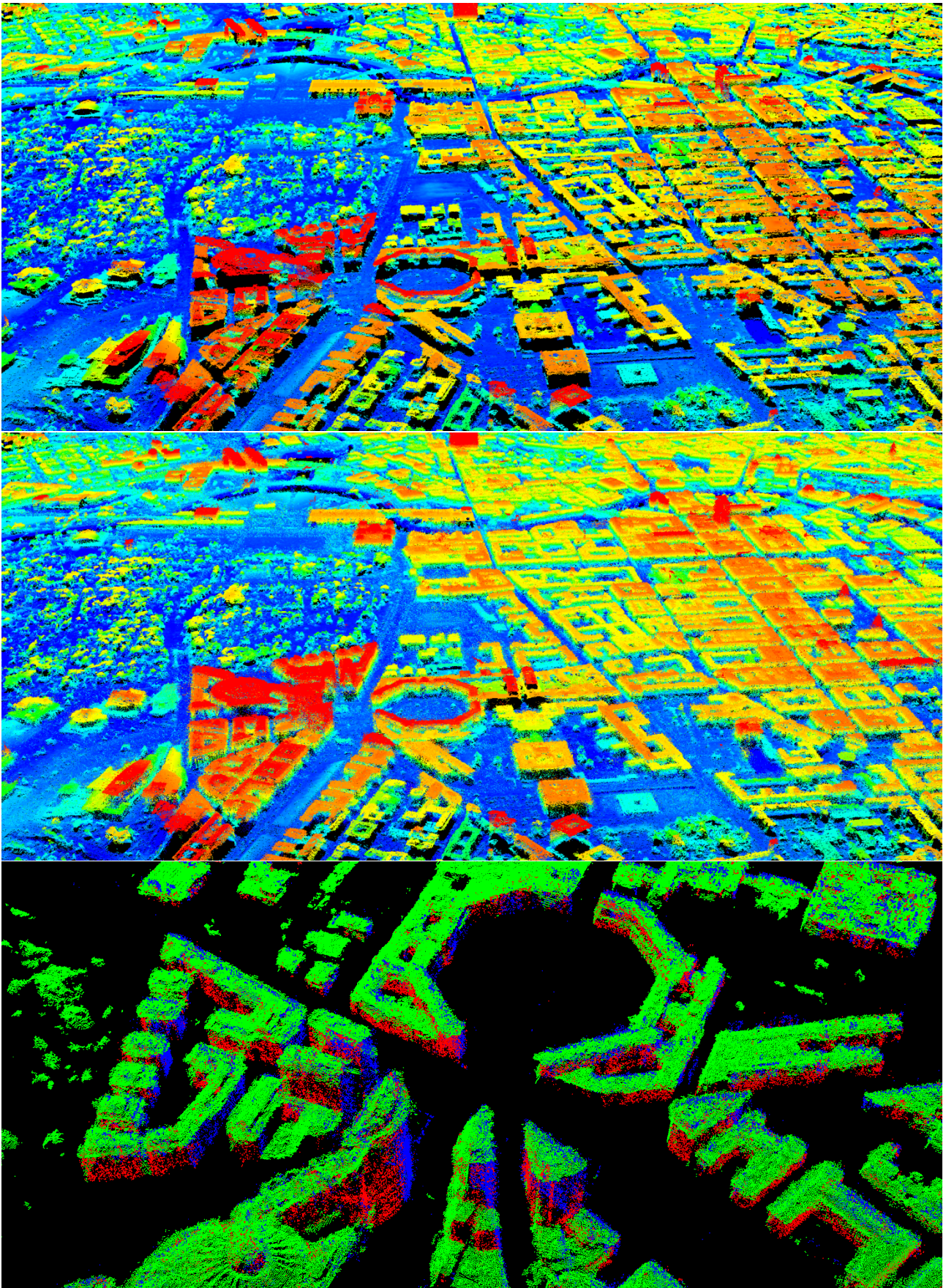
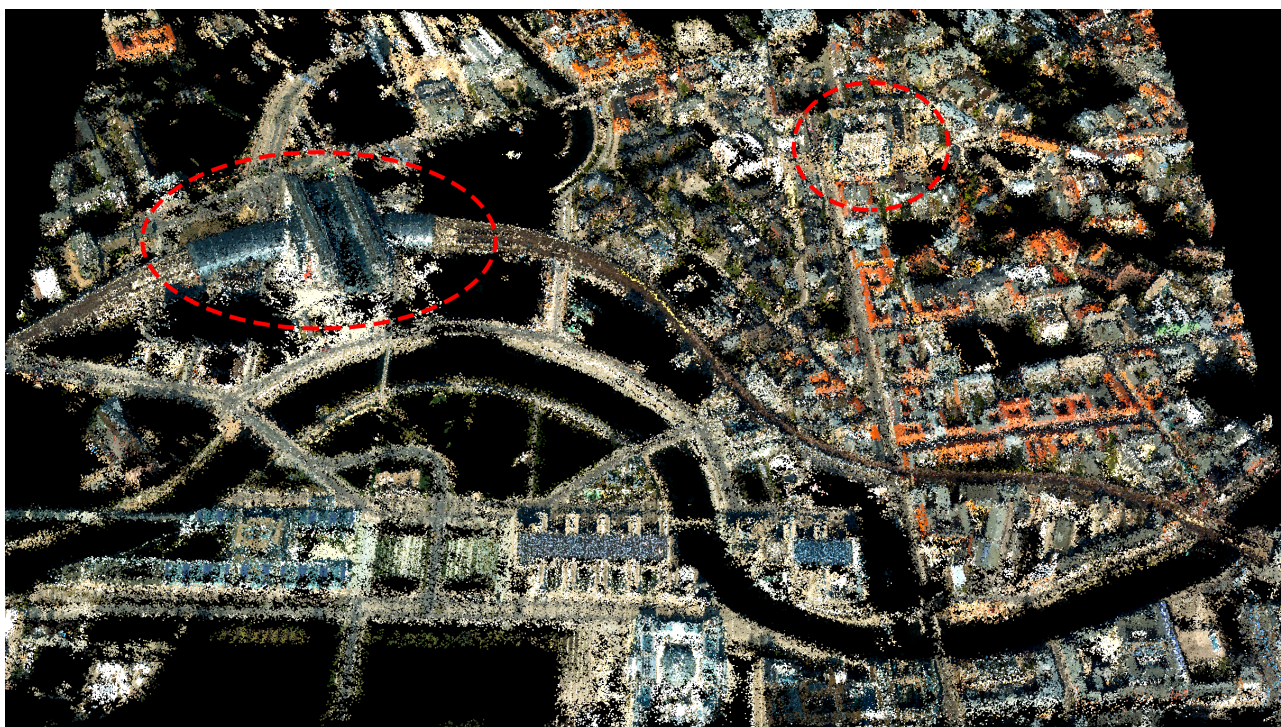
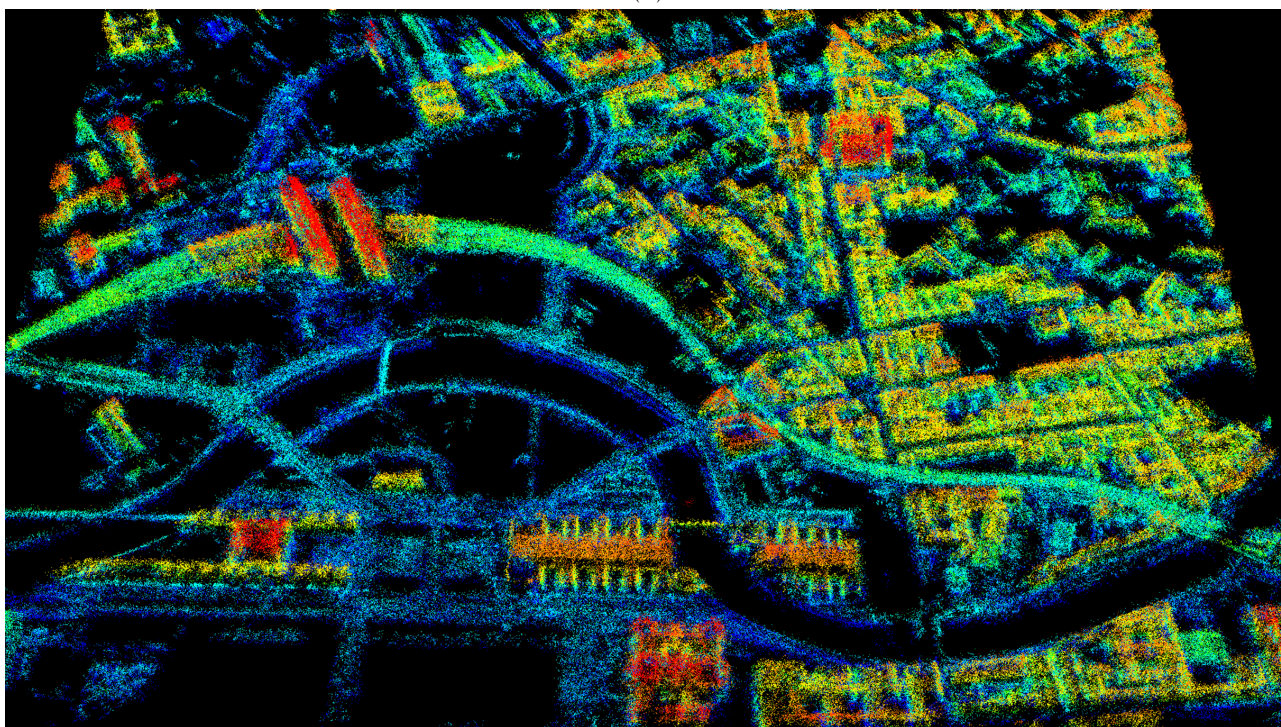


Fig. 4.16. Top: optical point cloud reconstructed using standard structure from motion algorithm with colour representing the height, middle: the fused point cloud combining the optical one and two TomoSAR point clouds from ascending and descending viewing angle, and bottom: a close up view of co-registered point cloud in Berlin Potsdamer Platz, where green, red, blue represent the points from optical, ascending and descending TomoSAR point cloud, respectively. The ground points were removed for a better visualization.



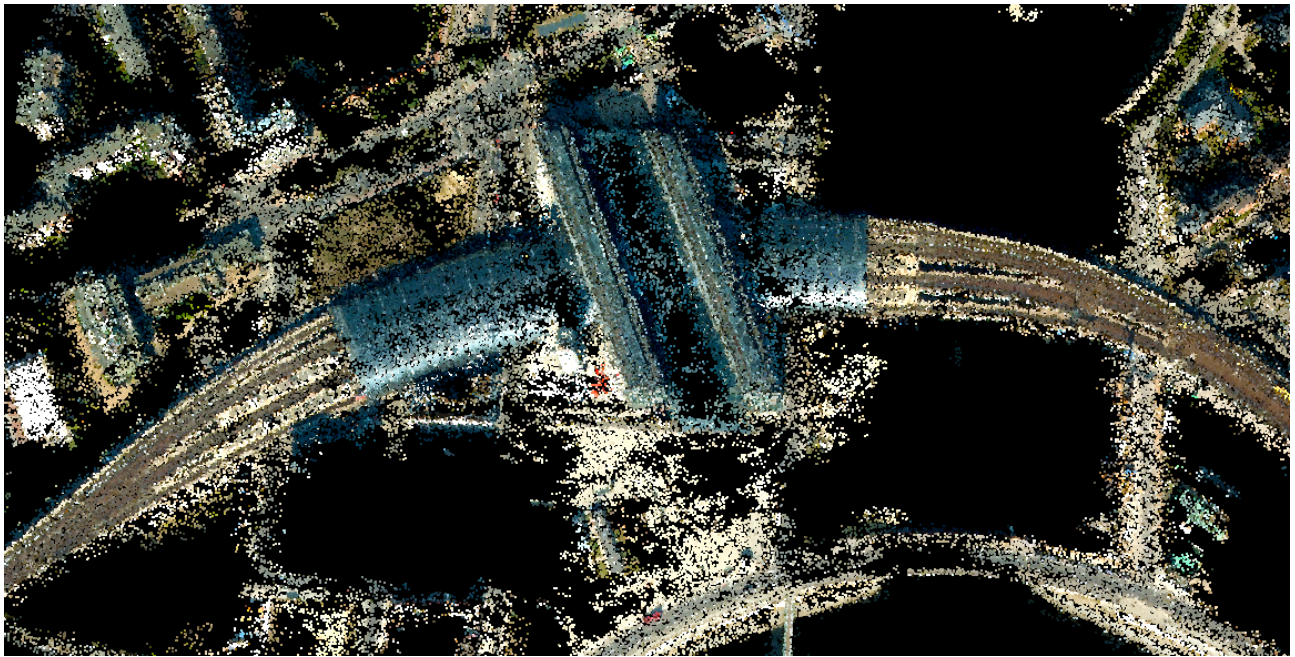


(a)



(b)





(c)



(d)

Fig. 4.17. (a) The first spaceborne meter-resolution 3-D TomoSAR point cloud of urban area textured with colour from optical image, (b) the corresponding TomoSAR point cloud colour-coded by height, (c) and (d) zoom in to the Berlin central station and Universitätsmedizin Berlin marked in dashed red ellipses in (a), respectively.

### 4.3.3 Practical Demonstration

#### *InSAR railway monitoring*



If classification is also performed on the optical image, the semantic labels can be transferred to the InSAR point cloud for further analysis. The details of the image classification are not presented here, as this is not the focus of this thesis. One can use any classification method. For this study, the Bag of Word (BoW) classification method (Csurka et al., 2004) was employed for classifying the railway and river class of an area near Berlin central station for the purpose of railway and bridge monitoring. The following figure shows the classification map where the railway class and river class are labelled in green and red, respectively.

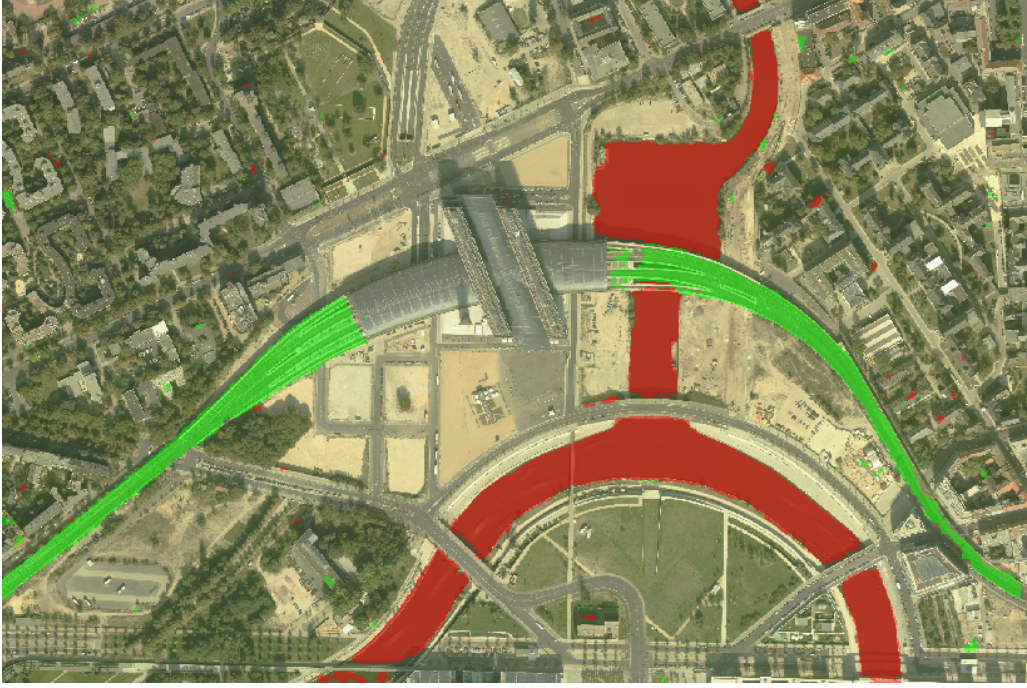


Fig. 4.18. River (red) and railway (green) classified using the BoW method.

Using the semantic label, the corresponding TomoSAR points can be extracted. Figure 4.19(a) shows the extracted continuous railway points overlaid on the optical image. The colour shows the amplitude of seasonal motion caused by thermal dilation. It can be easily observed that the railway deformation experiences certain discontinuity. We suspect it is because of the mechanical clearance between different railway segments. Therefore, we would like to detect and monitor these gaps.

However, the raw TomoSAR estimates are relatively noisy due to the relative low SNR of some scatterers. To filter out the inaccurate estimates, a special prior was employed, i.e. the thermal dilation of a steel beam is linearly proportional to its length at a first approximation (Kerr, 1978). That is to say, the railway deformation estimates are piecewise linear, and the scatterers on the same cross-section of the railway undergo similar deformation. Therefore, the deformation estimates can be filtered by minimizing its total generalized variation (with second order derivative which favours piecewise linear prior):

$$\hat{\mathbf{f}} = \arg \min_{\mathbf{f}} \left\{ \frac{1}{2} \|\mathbf{f} - \mathbf{y}\|_2^2 + \lambda \|\Delta \mathbf{f}\|_1 \right\} \quad (4.4)$$

where  $\mathbf{y}$  is the deformation estimates along the 1-D railway direction (i.e. stretch the railway to a straight line),  $\mathbf{f}$  is the noise-free version of  $\mathbf{y}$ , and  $\Delta \mathbf{f}$  is the second order derivative of  $\mathbf{f}$ . As shown in literatures of total generalized variation (Bredies et al., 2010; Knoll et al., 2011), the  $L_1$  norm of second order derivative is convex and lower semi-continuous, one can solve it using convex optimization solvers.

The filtered deformation estimates are shown in Figure 4.19(b), in parallel with the unfiltered estimates. The piecewise linear deformation signal is well revealed. To have a quan-

titative comparison, the upper subfigure of Figure 4.20 plots the original (in blue) and the denoised (in red) deformation estimates as a function of the 1-D railway direction. The original estimates have a standard deviation about 1 - 2 mm which concealed some edges where the magnitude of deformation is low. The filtering successfully reconstructed these edges.

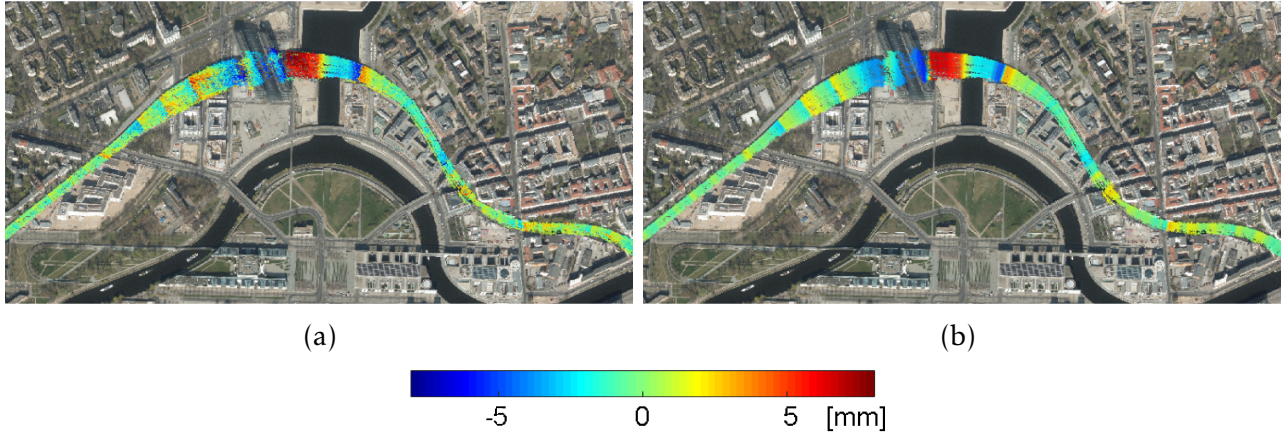


Fig. 4.19. (a) Railway points extracted from the original TomoSAR point cloud, and (b) the railway points filtered using total generalized variation. The colour shows the amplitude of seasonal motion due to the thermal expansion of the railway.

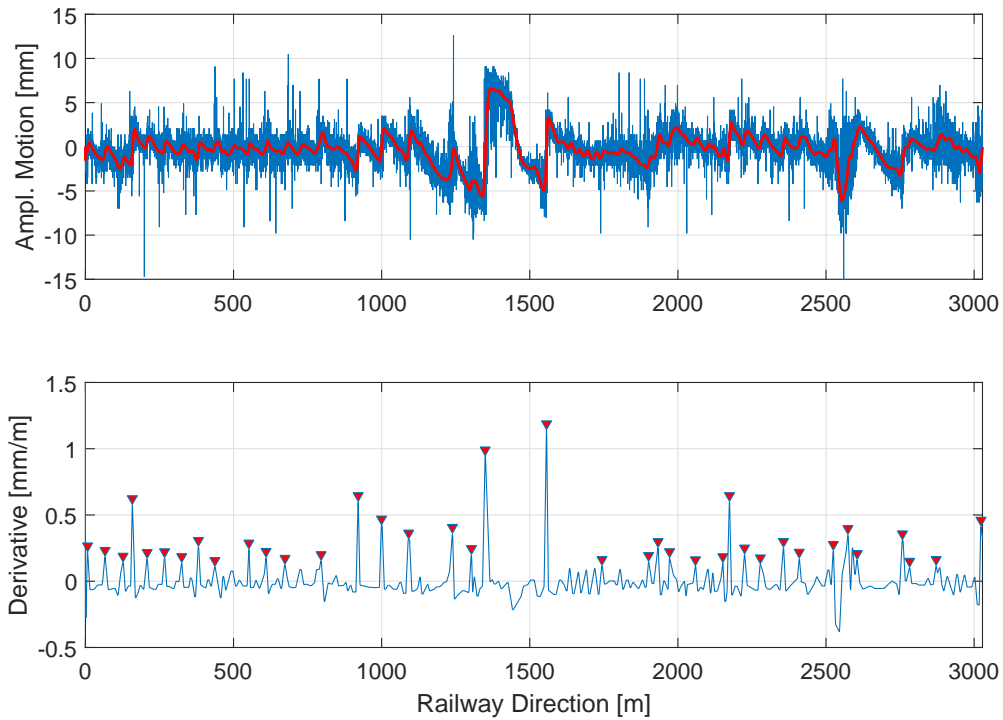


Fig. 4.20. Upper: the original and the filtered estimates of amplitude of seasonal deformation in blue and red, respectively, and lower: peaks detected in the derivative of the denoised deformation function along the railway direction.

By detecting the peaks in the derivative of the deformation function, the discontinuities in the deformation signal can be detected. Constraint was put on the minimum distance between two peaks. The detected peaks in the deformation's derivative can be seen in the lower subfigure of Figure 4.20. The positions of the peaks on the railway are shown as the green dots in Figure 4.21. Each green dot represents the midpoint of the cross-section. In the lower subfigure of 4.21, we provide the close up view of the two joints in the optical image. After carefully inspecting the optical images, we found that the several green dots close to the central station are indeed physical railway joints, while no physical mechanical joint is observed for the others green dots.





Fig. 4.21. Upper: the midpoint of the detected railway joint cross-section marked in green, and lower: close up view of the railway joint. The background optical image has a ground spacing of 20cm.

## 4.4 Robust Estimators for Multipass SAR Interferometry

Robust estimation has not been systematically addressed for multipass InSAR techniques. Appendix D and E address the robust estimation of two fundamental problems in the multipass InSAR algorithms. They are:

- ◇ covariance matrix estimation for non-Gaussian and nonstationary samples
- ◇ phase history parameters estimation for observations with large unmodelled phase errors.

The integrated new framework is called *robust InSAR optimization* (RIO). Table 4 summarizes the comparison of the state-of-the-art and RIO.

### 4.4.1 Robust Covariance Matrix Estimators

#### Non-Gaussian samples

The MLE for stationary CCG samples is the sample covariance matrix (2.12). RIO robustifies it using the M-estimator of the covariance which is basically an iterative reweighted sample

Table 4. Comparison of RIO framework with the state-of-the-art

	State-of-the-art	RIO
	Covariance estimation	
Stationary DS	Sample covariance matrix: $\hat{\mathbf{C}}_{MLE}$	M-estimator: $\hat{\mathbf{C}}_{M-est}$
Nonstationary DS		Rank M-estimator: $\hat{\mathbf{C}}_{RME}$
	Phase history parameters estimation	
PS	Periodogram: $\hat{\theta}_{MLE}$	M-estimator: $\hat{\theta}_{M-est}$
Stationary DS	Periodogram on samples mean: $\hat{\theta}_{MLE}$	M-estimator on samples mean: $\hat{\theta}_{M-est}(\hat{\mathbf{C}}_{M-est})$
Nonstationary DS	—	M-estimator pixel-wise inversion: $\hat{\theta}_{M-est}(\hat{\mathbf{C}}_{RME})$

covariance matrix (Ollila and Koivunen, 2003a; Zoubir et al., 2012):

$$\hat{\mathbf{C}}_{k+1} = \frac{1}{M} \sum_{m=1}^M w(\mathbf{g}_m^H \hat{\mathbf{C}}_k^{-1} \mathbf{g}_m) \mathbf{g}_m \mathbf{g}_m^H \quad (4.5)$$

where  $m$  and  $k$  are the sample index and the iteration index, respectively, and  $w(x)$  is a weighting function of the whitened squared residual  $\mathbf{g}_m^H \hat{\mathbf{C}}_k^{-1} \mathbf{g}_m$ . The weighting function down-weights highly deviated samples whose whitened residual is large in magnitude.

To derive a meaningful weighting function for some heavily tailed distributions, RIO models the non-Gaussian SAR pixel as a complex t-distribution (Tan, 1973; Kotz and Nadarajah, 2004):

$$f_{\mathbf{g}}(\mathbf{g}) = \frac{\Gamma(\nu + N)}{(\pi\nu)^N \Gamma(\nu) \det(\mathbf{C})^N} \left(1 + \frac{1}{\nu} \mathbf{g}^H \mathbf{C}^{-1} \mathbf{g}\right)^{-(\nu+N)} \quad (4.6)$$

where  $\Gamma(\cdot)$  is the Gamma function, and  $\nu \in \mathbb{R}^+$  is the DoF. The complex t-distribution is a realistic model for multipass SAR observations, as the DoF allows a quantitative definition of its deviation from CCG. This distribution approaches CCG as the DoF approaches  $+\infty$ , and becomes more heavily tailed as it decreases toward zero. In practical terms, this provides flexibility for handling different levels of outliers. The corresponding weighting function is derived according to equation (2.19):

$$w(x) = \frac{2N + \nu}{\nu + 2x} \quad (4.7)$$

For a good safeguard against outliers, a small value of  $\nu$  ( $<5$ ) is preferred (Ollila and Koivunen, 2003b). For the best safeguard against outliers, one can choose  $\nu = 1$ . This particular complex t-distribution with  $\nu = 1$  is also known as the complex Cauchy distribution, which has been thoroughly investigated in many studies (Feller, 1971). Besides, (Schmitt and Stilla, 2014) chose  $\nu = 2.5 - 5$  for centimeter resolution millimeter-wave SAR data of urban areas. This implies that the DoF is dependent on the object as well as the sensor. The simulation of different DoF is presented later in "quantitative comparison".

Equation (4.5) is solved iteratively. An approximation to drop the iteration is the sign covariance matrix (SCM) (Visuri et al., 2000; Croux et al., 2002). Extend it to complex number,



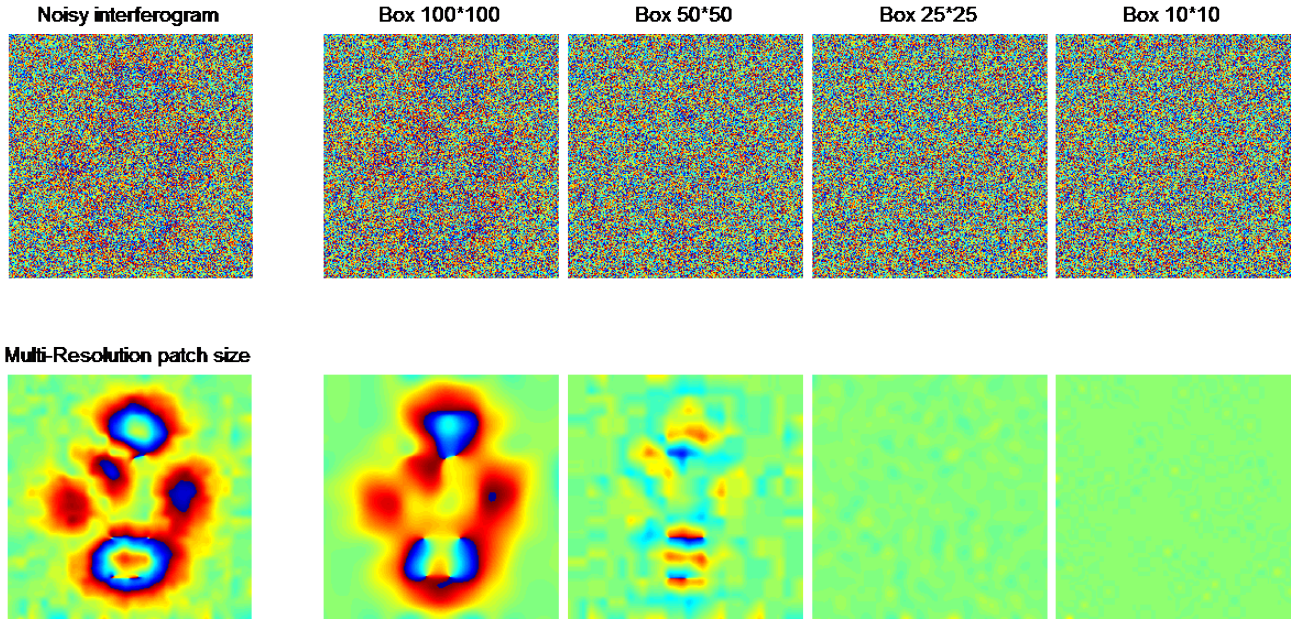


Fig. 4.22. Left: input noisy interferogram (upper), and the final estimated fringe (lower), and right: the input noisy interferogram (upper row) and estimated fringes (lower row) at each resolution level. The estimated fringes at each resolution level are subtracted from the noisy interferogram. The residual phase is taken as the input for the next resolution level with smaller window size.

it is

$$\hat{\mathbf{C}}_{SCM} = \frac{N}{M} \bar{I} \sum_{m=1}^M \|\mathbf{g}_m\|^{-2} \mathbf{g}_m \mathbf{g}_m^H \quad (4.8)$$

### Non-Gaussian and nonstationary samples

To complete the theory, we must consider the situation where samples are neither complex Gaussian nor stationary. The nonstationarity we consider is the uncompensated deterministic part of the interferometric phase, e.g. topography, motion etc. In this case, a joint estimation of the expected interferometric phase and the covariance (De Zan and Rocca, 2005) is required, i.e.,

$$\hat{\mathbf{C}}_{MLE} = \frac{1}{M} \sum_{m=1}^M \hat{\Phi}_m^H \mathbf{g}_m \mathbf{g}_m^H \hat{\Phi}_m \quad (4.9)$$

where  $\hat{\Phi}_m$  is the complex diagonal matrix containing the estimated phase values of  $\mathbf{g}_m$ . Usually, the estimation of the interferometric phase is performed by spatial filtering. Its performance greatly depends on the filter itself, which can be very challenging in urban areas.

In Appendix E, a multi-resolution defringe algorithm was developed to tackle this challenge. Figure 4.22 is a demonstration of the defringe algorithm. The left two figures are the simulated interferogram with the coherence of 30% of the pixels equal 0.3 and the rest equal to 0 (up), and the final estimated interferometric phase (bottom), respectively. The right eight subfigures show the input and estimated interferometric phase at each resolution level. Starting from the largest patch size, the fringe frequency is estimated using only the coherent DSs in a patch. The interferogram is then demodulated, and passed to the next resolution level. In the test, four resolution levels were set: 100×100, 50×50, 25×25, and 10×10 pixels.

Nevertheless, the phase estimation requires additional effort, and poor estimates significantly affect the covariance matrix estimation. Therefore, a new covariance estimator — *Rank M-estimator* (RME) — for complex multivariate is proposed, which is invariant to the

multiplicative phase signals. The RME is derived by replacing the multivariate  $\mathbf{g}$  with its rank  $\mathbf{r}$  in equation (4.5):

$$\hat{\mathbf{C}}_{RME,k+1} = \frac{1}{M} \sum_{m=1}^M w(x_m(\hat{\mathbf{C}}_{RME,k})) \hat{\mathbf{r}}_m \hat{\mathbf{r}}_m^H \quad (4.10)$$

The complex rank vector  $\mathbf{r}$ , being analogous to its real number version (Croux et al., 2002), is defined as follows:

$$\hat{\mathbf{r}}_m = \frac{1}{J} \sum_{j=1}^J \frac{\mathbf{g}_m \odot \mathbf{g}_j^*}{\|\mathbf{g}_m \odot \mathbf{g}_j^*\|} \quad (4.11)$$

where  $\mathbf{g}_j$  is the direct neighbourhood of  $\mathbf{g}_m$ , and  $\odot$  denotes the element-wise multiplication. The subtraction of neighbouring samples in the original definition is replaced by multiplication of the complex conjugate. The deterministic phase of  $\mathbf{g}_m$  is reduced through this operation. The RME is a fourth-order descriptor of the sample statistics. It can be proven that the element-wise square root of  $|\hat{\mathbf{C}}_{RME}|$  approaches  $|\hat{\mathbf{C}}_{MLE}|$  asymptotically under CCG distribution when calculating the rank using one neighbourhood sample. Due to the multiplication operation, the variance of the square-root  $|\hat{\mathbf{C}}_{RME}|$  estimate is doubled under a CCG distribution compared to  $|\hat{\mathbf{C}}_{MLE}|$ . Therefore, the number of samples should be doubled when using RME in order to keep a comparable variance as MLE. For an effective phase cancellation in equation (4.11), the neighbourhood should be kept small. We usually restrict  $\mathbf{g}_j$  to be the only four adjacent pixels of  $\mathbf{g}_m$ . Hereafter, we assume that the element-wise square root on  $|\hat{\mathbf{C}}_{RME}|$  has been performed.

### Qualitative comparison

We compared the original MLE, the M-Estimator with complex t-distribution weighting, and the RME under three different scenarios: (1) multivariate CCG, (2) multivariate complex t-distribution with one DoF, and (3) nonstationary multivariate complex t-distribution with one DoF. The data are simulated to have ten acquisitions, with a neighbourhood of 1000 samples. Each ten-variate sample vector is simulated by de-whitening a ten-variate vector of zero-mean i.i.d. CCG distribution (for scenario 1) or complex t-distribution (for scenarios 2 and 3) with a predefined coherence matrix. In the last scenario, linear fringes with ten different fringe frequencies randomly picked within  $[0 \pi/10]$  were added to the phases of the ten acquisitions.

In addition, two different coherence matrices were predefined; one exponentially decaying and the other with a constant 0.5 coherence between acquisitions. The results are shown in Figure 4.23 where the left subfigure corresponds to the exponential coherence matrix, and the right corresponds to the constant matrix. Each row represents one of the aforementioned three scenarios, and each column represents one of the three covariance estimators. The top left subplots in both subfigures are regarded as their reference coherence matrices, respectively, because the MLE is the optimal estimator under CCG and is asymptotically unbiased.

All three estimators can preserve the correct shape of the covariance matrix under CCG. The MLE fails in the second scenario, where the samples are contaminated by outliers. The coherence is usually overestimated because of the large amplitude of the outliers. Nevertheless, the M-estimator and RME remain as correct estimations. In particular, this M-estimator is the MLE for the simulated data. Finally, the M-estimator is not capable of dealing with nonstationary samples. Heavy underestimation occurs because of the summation of the complex numbers with nonstationary phases. The degree of underestimation depends on the local fringe frequency and the spatial extend of the samples. The larger the fringe frequency and

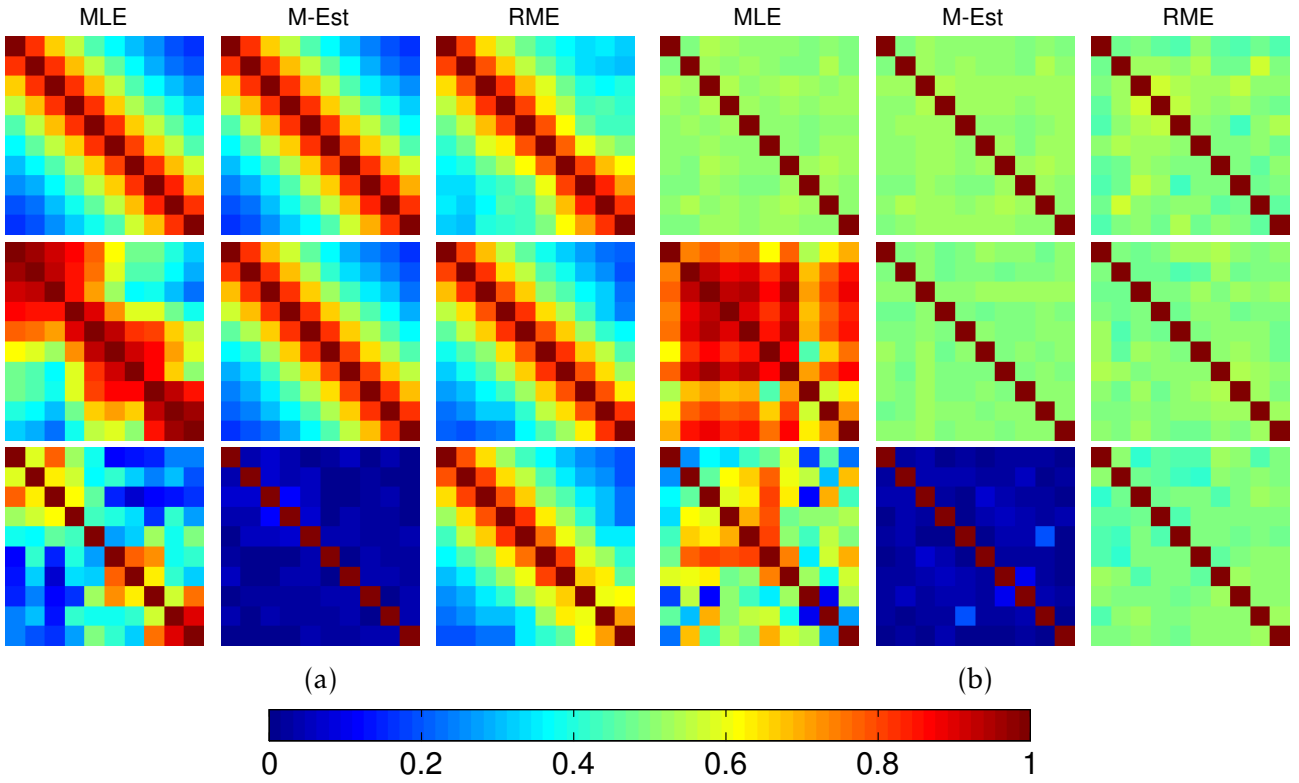


Fig. 4.23. Comparison of three covariance matrix estimators under three different observation cases; 1st row: complex circular Gaussian, 2nd row: complex t-distribution with one degree of freedom, and 3rd row: nonstationary complex t-distribution with one degree of freedom. (a) and (b) show the results of the exponential and constant coherence matrices, respectively. In both (a) and (b), 1st column: MLE (under Gaussian), 2nd column: M-estimator with t-distribution weighting, and 3rd column: rank M-estimator with t-distribution weighting.

the spatial extend of samples, the severer the underestimation. The RME is mean invariant, and maintains good performance in all conditions.

$|\hat{\mathbf{C}}_{RME}|$  shows higher bias at low coherence (as observed in Figure 4.23(a)), and also experiences higher variance (as observed in Figure 4.23(b)), due to the multiplication of two samples. This can be solved by using more samples in the estimation. The efficiencies of the M-estimator and RME are always lower than 100% under CCG distribution. This is a trade-off between robustness and efficiency.

### Quantitative comparison

The abovementioned example demonstrates the effectiveness of the robust estimator at the worst case, i.e. complex Cauchy distribution whose variance is infinite. For a systematic analysis of the robust covariance estimator at different levels of outlier percentage, the DoF of the complex t-distribution was varied from 2 to 5. As it was mentioned before, complex t-distribution approaches complex Gaussian as the DoF increases. Therefore, the advantage of the robust estimator diminishes as the DoF increases. Figure 4.24 shows the variance (upper) and bias (lower) of coherence estimates. The left side is for data with complex t-distribution of DoF=2.5, and the right side is DoF=5. The advantage of robust estimator is much less at DoF=5 than DoF=2.5, because the distribution of the former is more Gaussian. In this experiment, the robust estimator always outperforms the Gaussian MLE, because the robust weighting function is t-distribution weight. Should some empirical robust function be employed, the Gaussian MLE will outperform the robust estimator at certain point as the DoF increases. Since we are particularly interested in scatterers of low coherence, the variance and bias as a function of the DoF for true coherence = 0.2 were also plotted in Figure 4.25. As we can see, the performance of the robust estimator is very stable for different level of outliers.



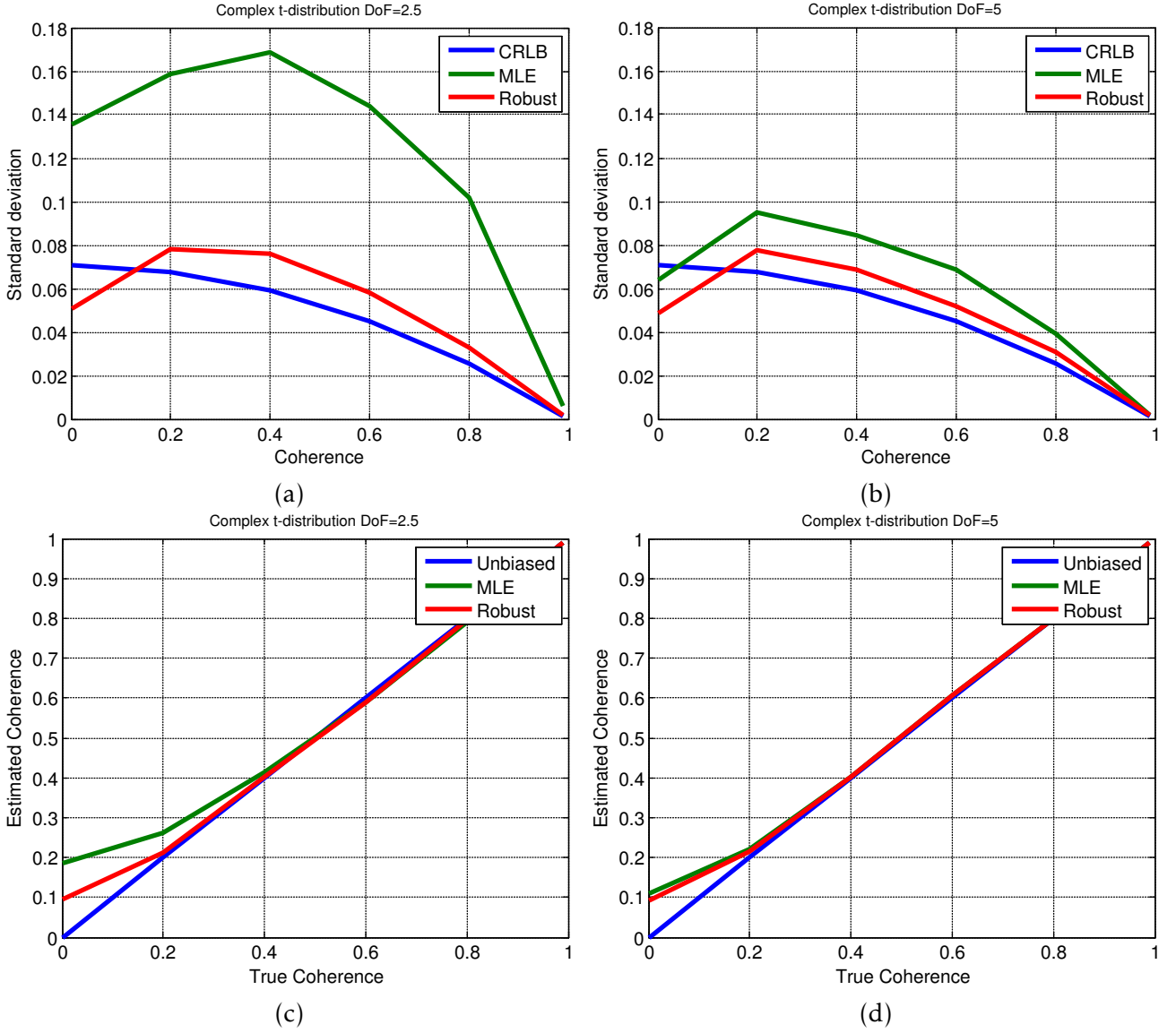


Fig. 4.24. (a) and (b): standard deviation of the coherence estimates of Gaussian MLE and the proposed robust estimator for complex t-distributed data with degree of freedom of 2.5 and 5, respectively, (c) and (d): bias of the coherence estimates. 100 looks is used in the coherence estimation.

#### 4.4.2 Generalized Rank Covariance Estimator

The abovementioned covariance RME is a robust estimator based on the first order phase derivative. It can be generalized to any order, so that the estimator will be invariant to any spatially continuous interferometric phase signal.

Consider a SLC observation  $g_j$  with image index  $j$ , its first order basic rank operator  $R^1[\cdot]$  is defined as the multiplication of itself with the complex conjugate of its neighbour pixel:

$$r_j = R^1[g_j] = g_j g_{j+1}^* \quad (4.12)$$

Assuming  $g_j$  being a Gaussian scatterer, and  $g_j$  being independent from  $g_{j-1}$ , the PDF of  $r$  is the product of two identical and independent CCG distribution. It is equivalent to the PDF of interferometric pixel with zero coherence, which was derived to be:

$$f(|r|, \phi) = \frac{2|r|}{\pi \bar{I}^2} K_0\left(\frac{2|r|}{\bar{I}}\right) \quad (4.13)$$

which does not depends on the phase  $\phi$  of  $r$ . Given a coherence of  $\gamma$  between two single look

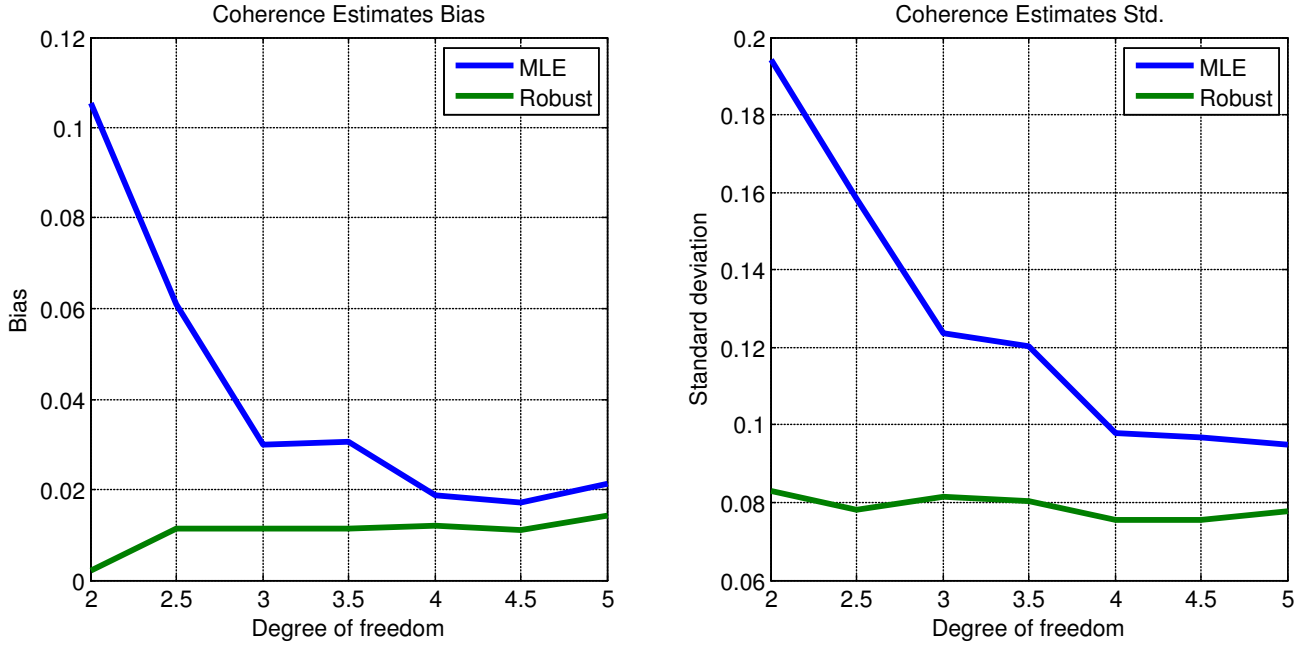


Fig. 4.25. Bias (left) and standard deviation (right) of the coherence estimates of Gaussian MLE and the proposed robust estimator as functions of degree of freedom of the complex t-distribution, for true coherence = 0.2. 100 looks is used in the coherence estimation.

observations, the second order moment of their first order basic rank, let's call it *first order rank coherence*, can be proofed to be  $\gamma^2$  by substituting equation (4.12) into the coherence equation (2.8).

Now the  $n$ -th order rank can be defined as the product of two  $(n-1)$ -th order rank pixel:

$$R^n[g_j] = R^{n-1}[g_j] R^{n-1}[g_{j+2^{n-1}}]^* \quad n \in \mathbb{N} \quad (4.14)$$

Please note that the multiplication is not performed with the pixel  $j+1$ , but  $j+2^{n-1}$ . This makes sure that an  $n$ th order rank pixel is the product of  $2^n$  independent SLC pixels. Therefore, the corresponding  $n$ th order rank coherence is

$$\gamma_n = \gamma^{2^n} \quad (4.15)$$

The variance of  $\gamma_n$  also increases accordingly, presumably by a factor of  $2^n$ . Therefore, the number of samples used in the estimation should also be increased by the same factor. The following figure compares the performance of the coherence estimation using different orders of rank. An interferogram is simulated with a constant coherence of 0.5 and the phase from the famous optical image of "Lena". The first subfigure shows the interferometric phase. The results from rank 0 to 2 are plotted in order. Particularly, rank 0 refers to the results using the original SLC pixels. It is obvious that heavy underestimation occurs if the original SLC is used. The degree of underestimation depends on the local fringe frequency. Rank 1 greatly reduced the underestimation. Rank 2 is able to eliminate all the underestimation. But the variance of the coherence estimates also increases as the rank increases.

To conclude, the generalized rank covariance estimator can be invariant to any differentiable spatial phase signal, if higher orders of rank are employed. However, more samples must be used for higher orders of rank, due to the fact that high variance is introduced by the rank operator. Rank 1 is a good balance between phase invariance and estimates variance.

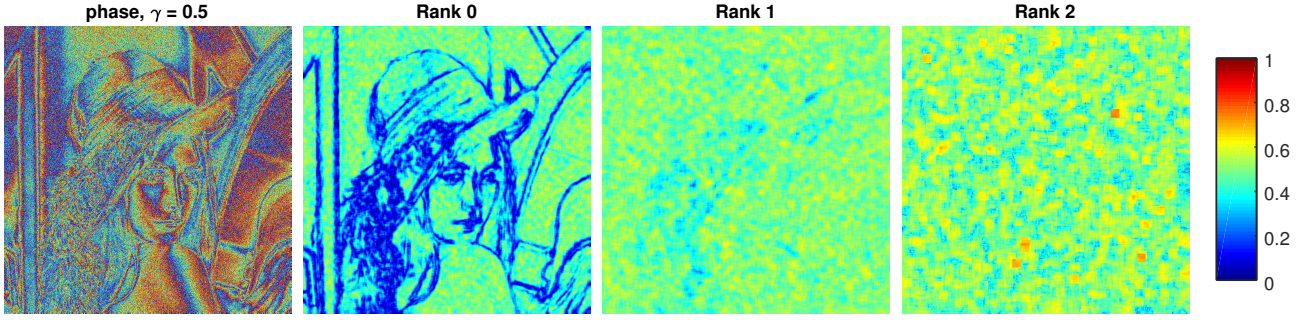


Fig. 4.26. Comparison of the coherence estimates using different orders of rank. The left subfigure shows the simulated interferometric phase with a constant coherence of 0.5. The other subfigures are the coherence estimates using rank 0 to 2. Rank 0 refers to using the original single look complex pixels. The colour bar indicates the coherence. Rank 0 heavily underestimate the coherence. Rank 1 greatly reduced the underestimation. Rank 2 is able to eliminate all the underestimation, but the variance of the coherence estimates also increases as the rank increases.

#### 4.4.3 Robust Phase History Parameters Estimators

This section describes how to robustly estimate the phase history parameters, i.e. elevation, and motion parameters, given a multipass InSAR observation  $\mathbf{g} \in \mathbb{C}^N$  with unmodelled phase, e.g. uncompensated atmospheric phase, unmodelled motion phase. The proposed methods were demonstrated using robust PS and DS phase history parameters estimators. This method can also be applied to other InSAR techniques, such as TomoSAR.

##### Robust PS Estimator

Recall equation (3.1) of PS MLE, which can be simplified as

$$\hat{\theta}_{MLE} = \arg \min_{\theta} \|\mathbf{g} - \bar{\mathbf{g}}(\theta)\|_2^2 \quad (4.16)$$

where  $\bar{\mathbf{g}}(\theta)$  is the modelled PS signal. Similar to the robust covariance estimator, it can be robustified by an M-estimator:

$$\hat{\theta}_{M-est} = \arg \min_{\theta} \sum_{i=1}^N \rho(\text{Re}[\varepsilon_i(\theta)]/\sigma_R) + \rho(\text{Im}[\varepsilon_i(\theta)]/\sigma_I) \quad (4.17)$$

where the residual  $\varepsilon_i(\theta) = g_i - \bar{g}_i(\theta)$ , and  $\text{Re}[\cdot]$ ,  $\text{Im}[\cdot]$  are the real and imaginary parts of a complex number, and  $\sigma_R$  and  $\sigma_I$  are the standard deviations of real and imaginary parts, respectively.

The  $\rho(x)$  function can be derived from the PDF of the contaminated distribution of  $\mathbf{g}$ , if it is known or can be modelled. However, it is usually unknown in practice. We shall use stable empirical functions instead, e.g. the Tukey biweight function introduced in section 2.4.3.

Equation (4.17) is solved iteratively, with  $\sigma_R$  and  $\sigma_I$  being updated at each iteration. Their initial values are obtained from the initial estimate  $\hat{\theta}_0$ . This is not critical for convex  $\rho(x)$ . However, a robust initial estimation is strongly recommended for nonconvex  $\rho(x)$ . The initial estimation need not to be efficient. To give one example, the least trimmed square that minimizes the sum of  $q$  smallest squared residuals (Rousseeuw, 1984) is as follows:

$$\hat{\theta}_0 = \arg \min_{\theta} \sum_{i=1}^q |\varepsilon_i(\theta)|^2 \quad (4.18)$$

The initial estimates of standard deviation, as well as those at each iteration, can be obtained using a robust estimator such as the median absolute deviation:

$$\hat{\sigma} = 1.483 \cdot \text{median}(|\varepsilon - \text{median}(\varepsilon)|) \quad (4.19)$$

The factor 1.483 is a normalization constant, which gives an expectation of unity for a normal distribution. With this combination, convergence is achieved in just a few iterations; therefore, the computational cost is only approximately three to four times more than that for the standard estimator. The same procedures can be easily extended to the LAMBDA estimator (Kampes and Hanssen, 2004) for PS phase parameter estimations.

### Robust DS Estimator

If stationarity is assumed for a DS and its neighbourhood, one can treat the DSs as a single PS by averaging them, as proposed in SqueeSAR. Then, the robustified DS estimator is identical to equation (4.17).

However, stationarity is not assumed in our considerations. The objective is a full inversion of individual single-look DS observation (DS pixel without averaging). The robustified estimator for DS phase history parameters is derived to be:

$$\hat{\theta} = \arg \min_{\theta} \left\{ \varepsilon^H(\theta) \mathbf{W}(\bar{\varepsilon}) \varepsilon(\theta) \right\} \quad (4.20)$$

The residuals  $\varepsilon(\theta)$  are whitened by a robust covariance matrix estimate, e.g.  $\hat{\mathbf{C}}_{RME}$ :

$$\varepsilon(\theta) = |\hat{\mathbf{C}}_{RME}|^{-0.5} \mathbf{\Phi}(\theta)^H \mathbf{g} \quad (4.21)$$

$\mathbf{W} \in \mathbb{R}^{N \times N}$  is the diagonal weighting matrix computed from each element of the expected residual  $\bar{\varepsilon}$ .

One should be aware that the marginal distribution of the phase of the residual is uniform, as it is in its whitened version  $\varepsilon(\theta)$ . The unexpected large phase error, presumably also uniformly distributed over time, applies no change to the DS observations statistically. In other words, the robust loss function is blind to phase contamination in single-look DS observations. Therefore, the correct weighting should be determined by the expected residuals  $\bar{\varepsilon}$  of the neighbourhood ensemble. The expected residuals should be calculated using a robust estimator:

$$\bar{\varepsilon} = \sum_{m=1}^M w^m \varepsilon^m / \sum_{m=1}^M w^m \quad (4.22)$$

where the superscript  $m$  denotes the sample number in the neighbourhood, and  $w^m$  is a robust weight.

To summarize, equation (4.20) is a joint estimation of the phase parameters of individual single-look DS observations in a neighbourhood. It is solved iteratively. Its computation should begin with initial estimates of each sample in the neighbourhood (assumed to be the same), which jointly determine the initial weighting matrix. The same weighting matrix is used to retrieve the parameters of each single-look DS in the neighbourhood, and is updated on the basis of all the estimates upon finishing one iteration. The pseudocode of the proposed DS estimator is as stated in the following table.

### Efficiencies of the Robust Estimators

The efficiencies of the proposed robust PS and DS estimators and their original MLEs are quantitatively assessed using Monte Carlo simulations. The sensor related parameters in the simulation were set to be similar to those of TS-X data, where the wavelength is 0.031 m, the range distance  $R = 700$  km, the acquisitions number  $N = 20$ , the spatial baselines  $\mathbf{b}_{\perp}$  were uniformly distributed between -100 m and +100 m, and the temporal baselines  $\mathbf{t}$  were evenly spaced from -1 year to +1 year with uniform distribution.

Table 5. Algorithm of the robust estimator for single-look DS observations

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```

Get initial estimate  $\hat{\theta}_0$ , which can be the same for all DS samples in the same neighbourhood
Calculate initial weighting matrix  $\mathbf{W}_0$ 
for iteration  $k = 1$  until convergence
    for DS sample  $m = 1$  to  $M$ 
        Estimate  $\hat{\theta}^m$  by solving equation (4.20) using  $\mathbf{W}_k$ 
    end
    Compute the expected residuals  $\bar{\epsilon}$  by a weighted average of  $\epsilon^m$  following equation (4.22).
    Compute  $\mathbf{W}_{k+1}$  based on  $\bar{\epsilon}$ 
end

```

---

The contamination-free PS and DS observations were simulated firstly. Then contamination is introduced in two ways: (1) random phase is added to eight (40%) of the 20 acquisitions. The phase was uniformly distributed between  $[-\pi, \pi]$  for both PS and DS. For DS, sufficient spatial samples were also simulated. The same phase is added to the samples of the same acquisition to simulate the temporally random but spatially stationary atmospheric phase, and (2) a complex t-distribution with one DoF, instead of CCG, was used in the DS simulation to simulate outliers in the DS samples. More details of the simulation can be found in Appendix E.

The proposed robust estimators and their original MLEs were applied to both the contamination-free and contaminated observations. The integral of Tukey's biweight function was chosen as the loss function for both robust PS and DS estimators. The parameter  $c$  was set to 4.586, and the Monte Carlo simulation was repeated 1000 times.

Figure 4.27 shows the standard deviation of the estimates of linear deformation rate with respect to different SNRs. The left and right subfigures correspond to the robust PS and DS estimators, respectively. The SNR of the DS refers to the correlated signal w.r.t. the decorrelated part, and can be directly related to the coherence by the formula  $\gamma = (1 + \text{SNR}^{-1})^{-1}$  (Zebker and Villasenor, 1992; Bamler and Hartl, 1998). In each subfigure, the green curves are the results of the proposed robust estimators, and the blue curves are the original MLEs; the dashed curves are the estimators performing on observations with CCG distribution noise, and the solid curves are results for contaminated data. The blue dashed curves in all the subfigures are the MLEs under CCG; hence, they are the optimal estimates.

Figure 4.27 conveys two important messages:

- (1) The results suggest that the robust estimators significantly outperform the original MLEs when large amount of the observations are contaminated. The robust estimators improve the efficiency by a factor of 7 to 35 (the squared ratio of the solid curves).
- (2) Still, the efficiency of the robust estimator under the nominal distribution (i.e. CCG) is close to that of the original MLE. The relative efficiency of the robust estimator and the MLE under a nominal distribution, defined as

$$\eta_{RE} = \frac{\sigma_{MLE}^2}{\sigma_{Robust}^2} \quad (4.23)$$

is approximately 70% for the proposed robust PS estimator and 80% for the DS estimator. Therefore, the robust estimators are also capable of handling uncontaminated

observations without too much loss of efficiency.

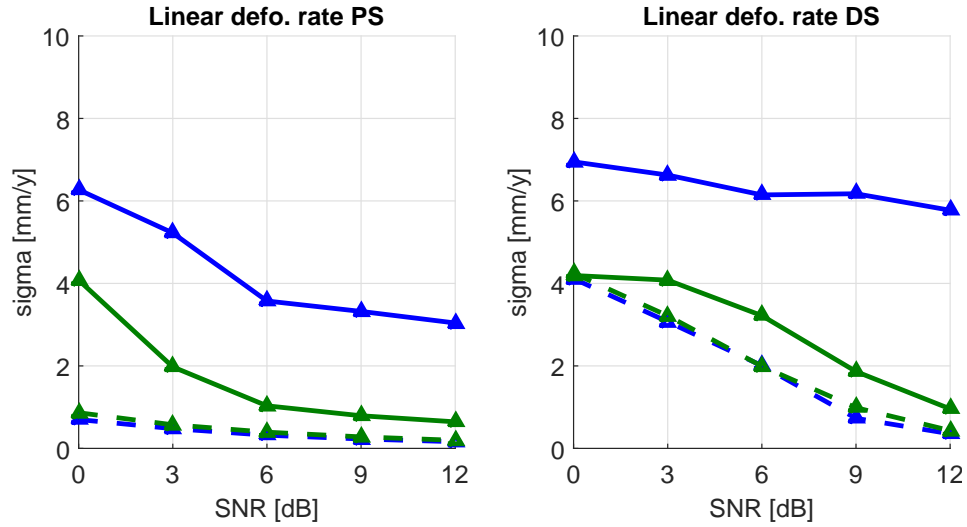


Fig. 4.27. Comparison of the standard deviation of the linear deformation rates estimated by the robust estimators and the ordinary MLEs. The left and right subfigures correspond to the PS and DS estimators, respectively. In both subfigure, the green curves are the results of the proposed robust estimators, and the blue curves are those of the original MLEs; the dashed curves are the estimators performing on contamination-free observations, and the solid curves are the results for contaminated data. The blue dashed curves in all the subfigures are the MLEs under a complex circular Gaussian distribution; hence, they are the optimal estimates. The robust estimators improve the efficiency by a factor of 7 to 35 for contaminated data (the squared ratio of the solid curves) while still keep a relative efficiency of 70% - 80% for contamination-free data (the squared ratio of the dashed curves).

#### 4.4.4 Practical Demonstration of the Robust InSAR Optimization

##### Against unmodelled phase

To demonstrate the robustness against large phase errors, an area in Las Vegas with significant non-linear motion was selected, in a stack of 30 TS-X high resolution spotlight images. The test area is marked by the red rectangle in Figure 4.28(a). The reference point marked by the red cross is a few hundred meters to the area. The correct deformation phase history of a sample pixel (marked by yellow cross) considering both linear and non-linear motion models is shown in Figure 4.28(b). As we can see, not only the motion is complex but also the magnitude of motion is very large. If only a linear motion model is considered, the unmodeled motion phase is equivalent to a large phase error. Non-robust estimators will give biased estimates.

Figure 4.29(a) compares the result of the proposed robust estimator and the ordinary MLE when only the linear model is considered. The robust estimates correctly reconstruct the subsiding bowl, whereas the MLE estimates are heavily biased by the unmodeled motion of the building roof. The advantage of the robust estimator is clearer in Figure 4.29(b) which is the bias of the estimates w.r.t. the correct linear deformation rate.

##### Against non-Gaussian scatterer

Figure 4.30 shows the difference between the linear deformation rate of the DSs estimated using the classical sample covariance matrix  $\hat{\mathbf{C}}_{MLE}$  (left column) and that using the proposed  $\hat{\mathbf{C}}_{RME}$  (right column). Identical samples were used for estimating these two covariance matrices. They were adaptively selected with the KS test using ten amplitude images, which were expected to introduce a non-negligible number of outliers in the selected samples because of the low detection rate from the low number of images. The same ordinary DS MLE was employed to estimate the linear deformation rate in both cases. Therefore, any improvement was solely credited to the use of a more robust covariance matrix estimate.



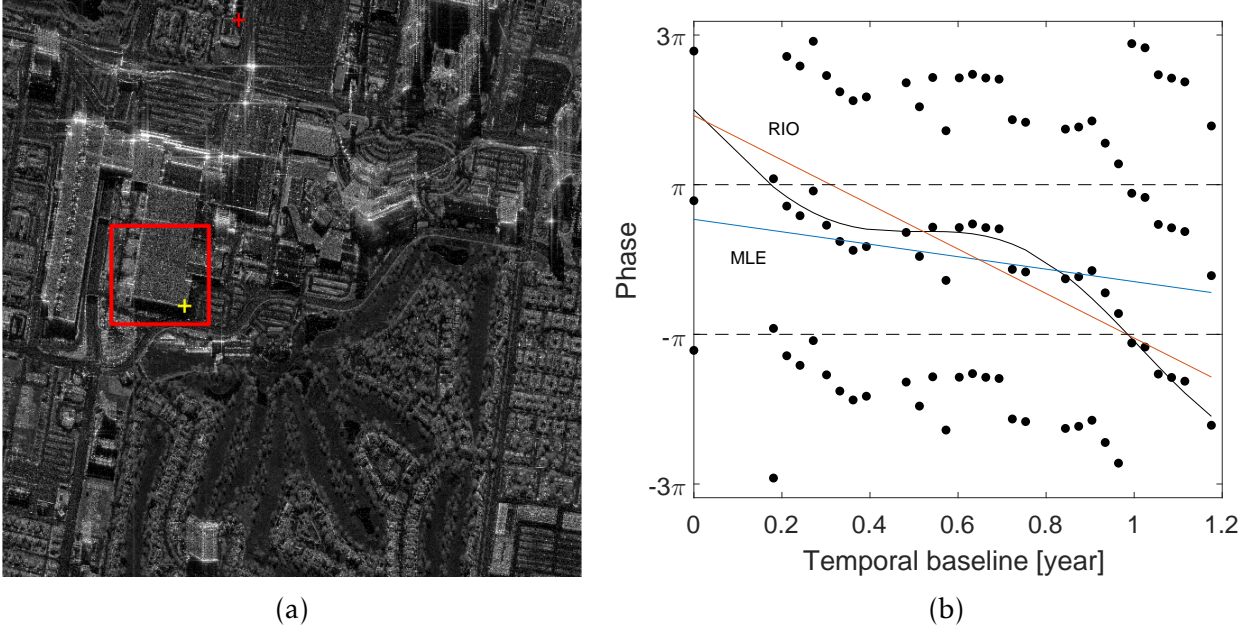


Fig. 4.28. (a) the test area with significant non-linear motion is marked by the red rectangle. The red cross is the reference point. (b) the deformation phase history of the pixel marked by yellow cross in (a), where the black dots are the wrapped deformation phases and its duplicates by adding and subtracting  $2\pi$ , the black curve is a non-linear fit to the phase history, and the red and blue lines are linear fittings using RIO and MLE, respectively.

The improvement is clearly demonstrated. The test area is mainly vegetation, except a road in the center which usually appears as a DS in the X-band images. Homogeneous deformation rates are expected, as the spans of both areas were roughly one hundred meters. However, many estimates appear in subfigure (a) as salt-and-pepper noise. We believe this is due to the low detection rate of the KS test, and the nonstationarity of the samples. In contrast, homogenous deformation rate of the road is shown in subfigure (b).

For further comparison, the histograms of the linear deformation rates of the road in Figure 4.30(a) and (b) were plotted in Figure 4.30(c) and (d) respectively. When using  $\hat{C}_{MLE}$ , many local peaks appear in the histogram which should not correspond to the deformation signal. With  $\hat{C}_{RME}$ , results are considerably more homogenous, and thus more reasonable.

### Big Area Demonstration

The RIO framework was applied on a stack of 33 TS-X high resolution spotlight images of the super volcano Campi Flegrei. Figure 4.31 shows the results of the linear deformation rate using the RIO framework, in which Tukey's biweight loss function is used for both robust PS and DS estimator. We specifically chose the SCM for the robust covariance estimate because of the absence of significant fringes in the interferogram.

RIO retrieved 15 times more scatterers than that using only PS. This number should also be valid for standard PS+DS methods, e.g. SqueeSAR. However, RIO greatly improves the overall robustness of the current multipass InSAR estimators. This can be seen in the small area comparison (Figure 4.32) of the two random areas marked by red rectangles in Figure 4.31, where RIO greatly reduced the salt and pepper noise which is probably caused by unmodeled phase and bad covariance estimation.

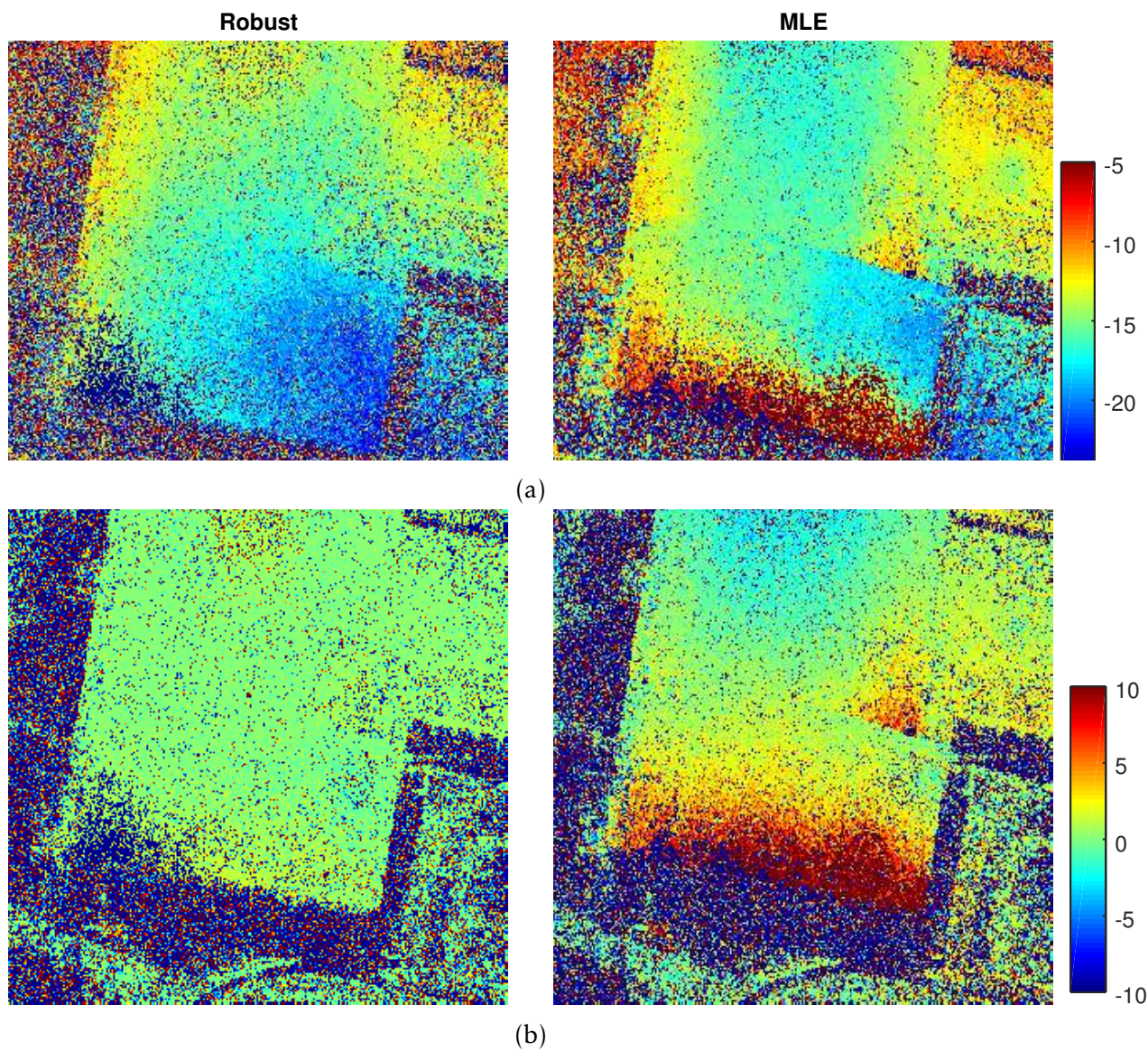


Fig. 4.29. (a) linear deformation rate estimates by robust estimator and ordinary MLE, and (b) the bias of the robust and non-robust estimates to the "ground truth" (linear deformation rate estimated considering the non-linear motion model). Units: [mm/year].



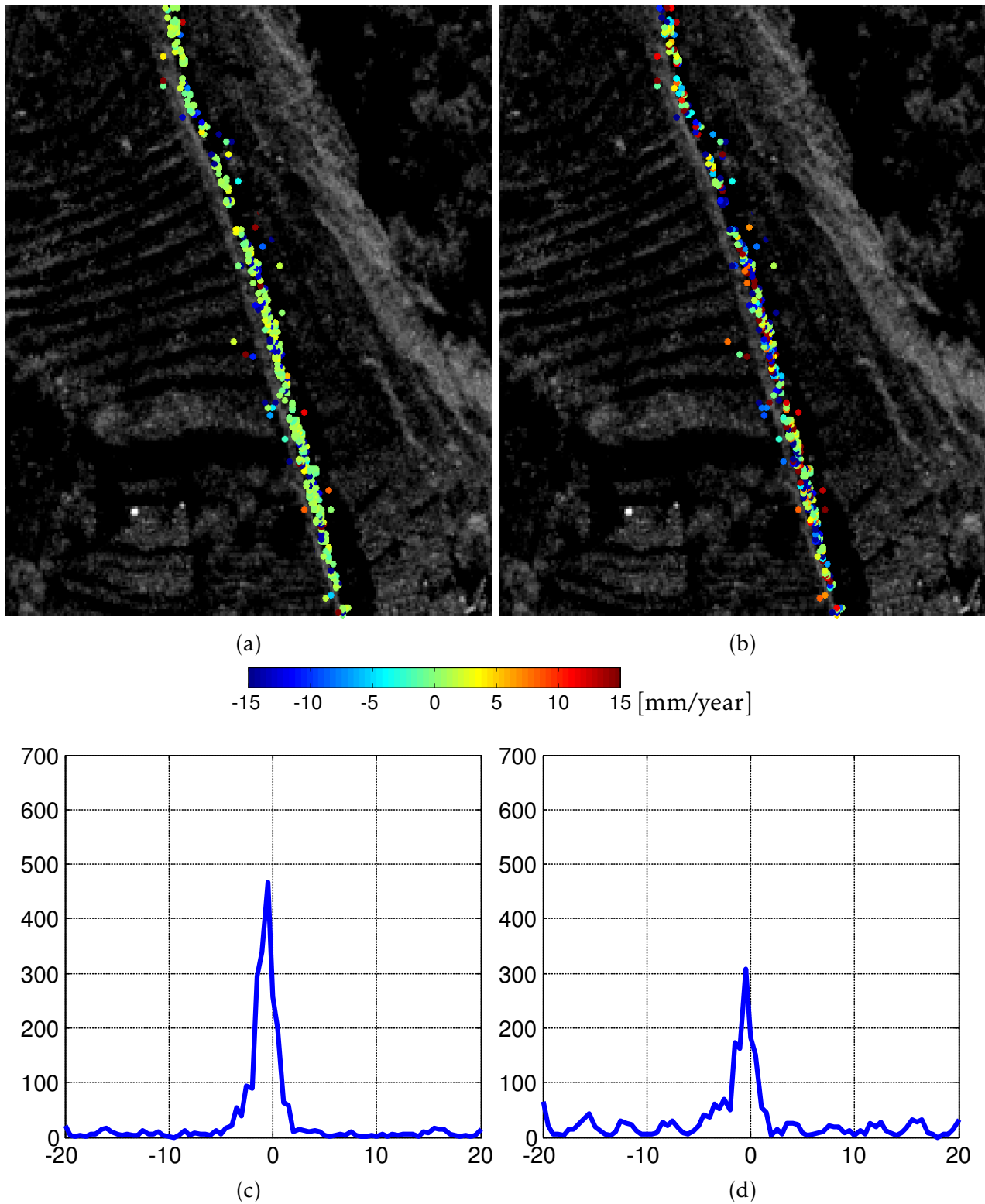


Fig. 4.30. (a) and (b) comparison of the linear deformation rate of the two test sites estimated using the proposed  $\hat{C}_{RME}$  and the ordinary  $\hat{C}_{MLE}$ , respectively. Identical samples were used for estimating both covariance matrices. They were adaptively selected (KS test) using ten images. This is expected to introduce a non-negligible number of outliers in the selected samples because of the low detection rate from the low number of images. Results (overlaid on the SAR intensity images) are specific to a local reference point. A homogenous deformation rate is expected because of the small size of the test area. The same ordinary DS MLE was employed for the linear deformation rate estimations in both cases. And (c) and (d) the corresponding histograms of the linear deformation rates in (a) and (b), respectively. In (d), many local peaks appear almost uniformly in the search range, most of which should not correspond to the deformation signals.



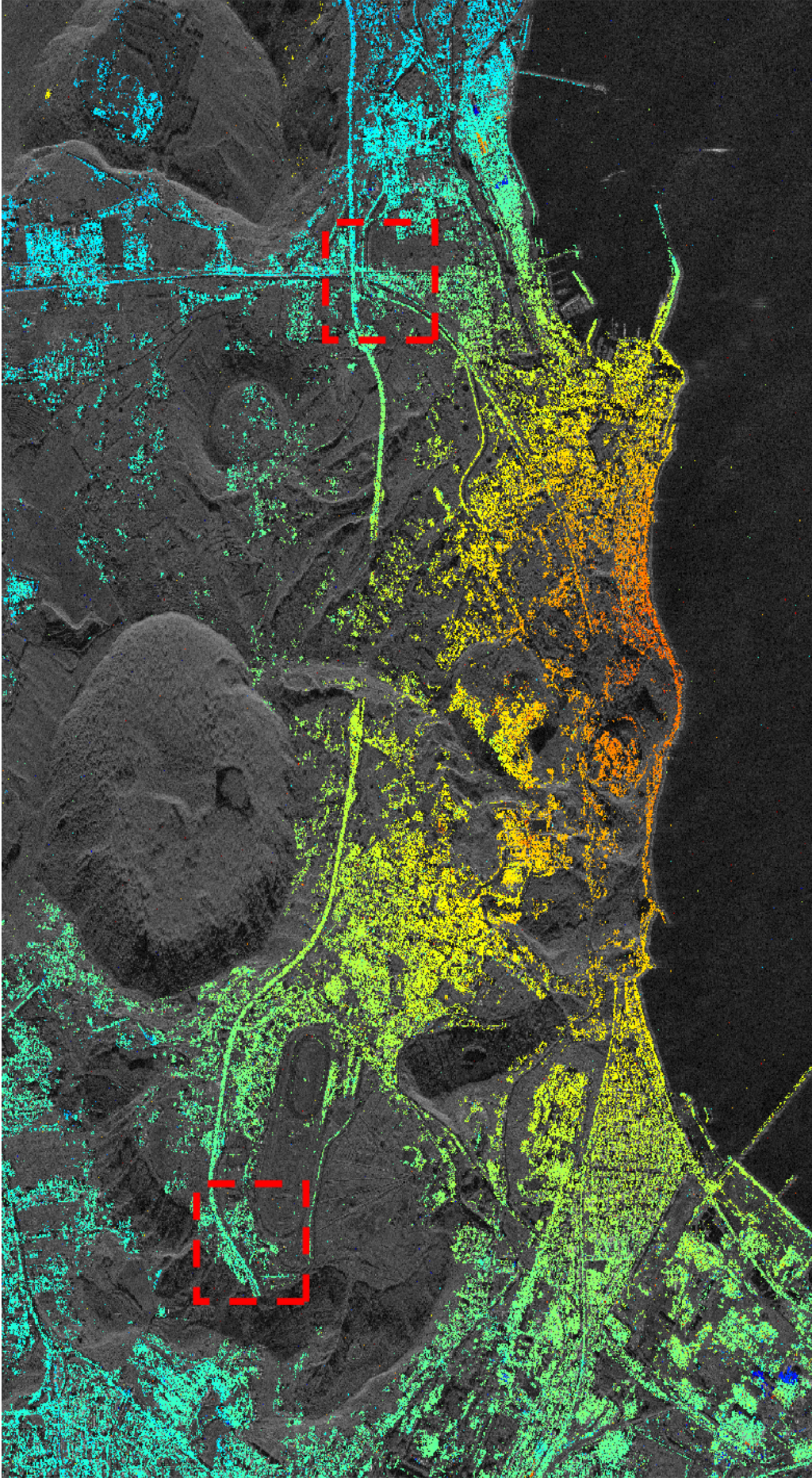


Fig. 4.31. Linear deformation rate of the super volcano Campi Flegrei, as retrieved using the proposed robust InSAR optimization framework that combines both PS and DS in a stack of 33 TS-X high-resolution spotlight images. Tukey's biweight function is set to be the robust loss function, and the sign covariance matrix is employed as the robust covariance matrix estimate. RIO retrieves 15 times more scatterers than those using only PS only. This number should also be valid for standard PS+DS methods, e.g. SqueeSAR. However, RIO greatly reduced the salt and pepper noise probably caused by unmodeled phase and bad covariance estimation. This can be seen in the small area comparison in which plots the results for the two areas marked by red rectangles.



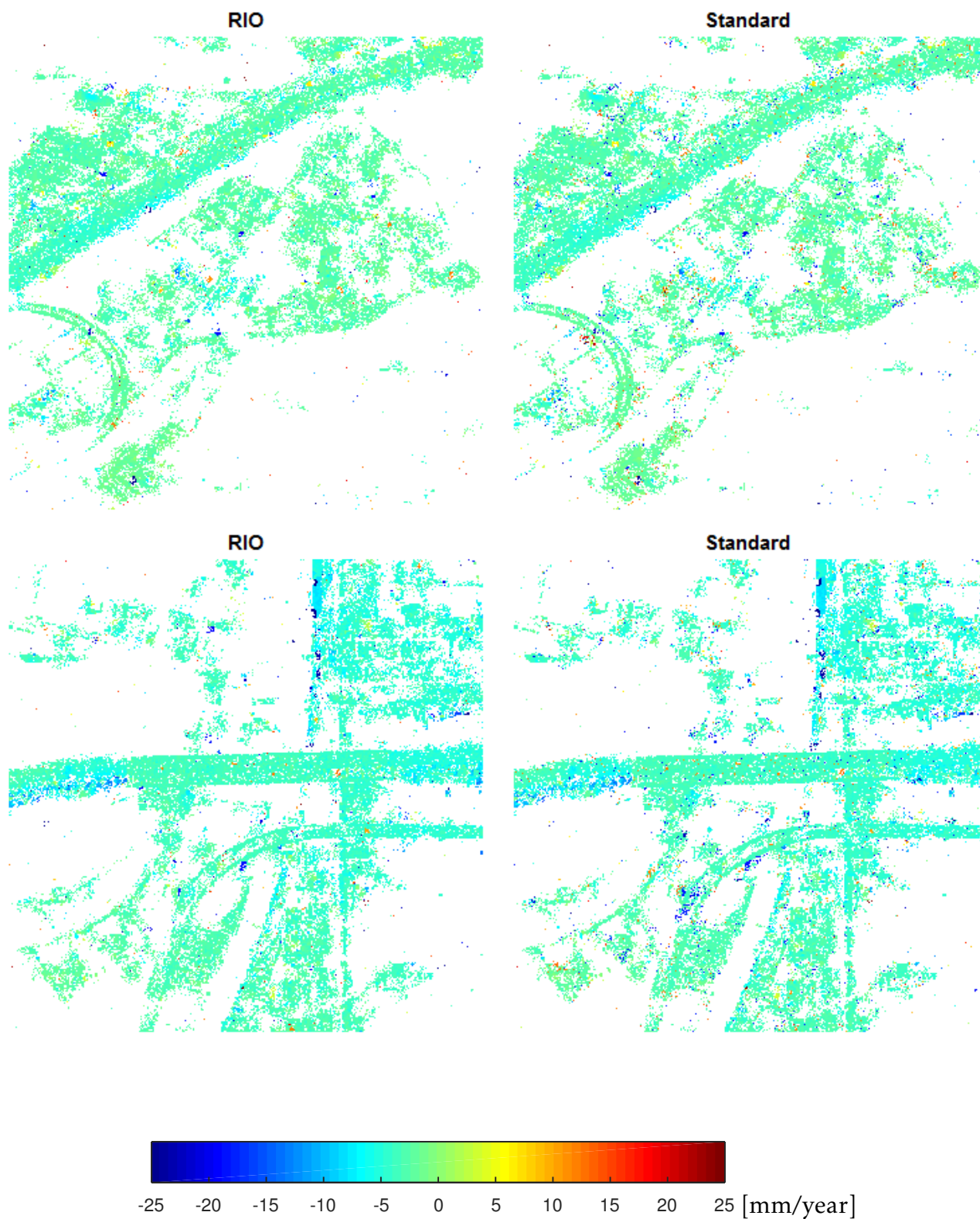


Fig. 4.32. Zoom in of the two areas marked by red rectangles in 4.31.

## 5 Conclusion and Outlook

### 5.1 Conclusion

**T**his dissertation is to develop advanced multipass InSAR algorithms, with the aim of computational efficiency, information fusion, contextual awareness, and statistical robustness. The objectives are fulfilled with the work presented in this thesis. The following contributions can be drawn.

- ◊ The high computational cost of tomographic SAR inversion compared to PSI requires an efficient approach for operational processing. The computational cost can be greatly reduced by
  - Pre-classifying single and double scatterers without performing TomoSAR inversion on every pixel. The classification should use simple quantities. We used pixel mean intensity, amplitude dispersion index, and the likelihood to single scatterer model for the pre-classification. The current implementation can achieve 70% double scatterer detection rate with 25% false alarm rate. The falsely detected double scatterer can be discriminated in the full tomographic SAR inversion afterwards.
  - using the estimates from PSI as a prior in the TomoSAR inversion of double scatterers.
  - and pass only the necessary pixels to the most expensive superresolution algorithms, e.g. SL1MMER. In our approach, only the single scatterers detected from the non-superresolution algorithm (e.g. SVD-Wiener) whose elevation estimates are less than 1.5 times of the Rayleigh resolution were passed.
 The computational cost of a large area can be reduced by a factor of 50.
- ◊ The interferometric phase often contains unmodeled contributions, e.g. unmodeled deformation and uncompensated atmospheric phase. And this is particularly true for very high resolution data of urban area, due to its complex motion and atmospheric phase with possibly high spatial frequency. Robust estimation is recommended for such case. The proposed robust estimator can improve the variance of the deformation estimates by a factor of 7 to 35, while still maintain a relative efficiency of 70% to 80% w.r.t. the MLE for uncontaminated data.
- ◊ The expected interferometric phase must be removed in prior of the covariance estimation. It is commonly called defringe, and commonly realized by spatial filtering. The filtering can be replaced by a simple subtraction of the phase of neighbouring pixel, i.e. multiply the complex conjugate of the neighbouring pixel. The augmented observation is called rank observation. The rank covariance estimator is invariant w.r.t. very large fringe frequency. The rank covariance can be generalized to any order, and be theoretically invariant to any fringe frequency.
- ◊ The performance of adaptive sample selection in SAR image stack decreases as the number of images decreases. Robust estimation of the covariance is necessary. The M-estimator of covariance provides a general framework. Among them, the non-iterative sign covariance matrix is shown to be a promising method for fast covariance estimation of low coherent signals.
- ◊ SAR images as well as InSAR estimate are difficult to interpret, especially for high resolution data of urban area. By geometrically fusing the optical image to the 3-D InSAR point cloud, we can intuitively understand the properties of the scatterers in visible spectrum. In this thesis, the first meter-resolution TomoSAR point cloud textured with optical colour was demonstrated.
- ◊ The current multipass InSAR techniques are very much pixel-based inversion and phase history analysis. By transferring the semantic classification label of the optical image to the InSAR point cloud, one can perform a systematic analysis of the InSAR point cloud

based on semantics, e.g. automatic railway monitoring.

## 5.2 Outlook

The future SAR and InSAR algorithms will become more intelligent and user friendly, which can be seen in the following aspects.

### Optical and SAR Image Joint Reconstruction

Some pioneer studies on joint pixel processing are emerging. For example, joint iso-height pixels TomoSAR (Zhu et al., 2015a,b; Ge et al., 2015), contextual phase unwrapping (Baselice et al., 2014), etc. Still, there is much information not fully exploited in SAR images. One obvious way is to use the rich spatial information in optical images to help the InSAR inversion. The following research directions are foreseen:

- ◇ Efficient and precise co-registration algorithm for very high resolution optical and SAR images in urban area, instead of full 3-D reconstruction from SAR and optical image introduced in this thesis.
- ◇ How to use the features in the optical image, e.g. the colour, classification labels, etc., to help the InSAR reconstruction. One way is to set different prior for different classes. Recall the TomoSAR MAP estimator in equation (3.19), better results can be achieved by introducing different prior covariance matrix  $C_{\gamma\gamma}$  for the prior of every class:

$$\hat{\gamma}_{MAP} = \min_{\gamma} \|\mathbf{g} - \mathbf{R}\gamma\|_2^2 + \lambda \|\gamma\|_{C_{\gamma\gamma}}^2 \quad (5.1)$$

Another direction is to find similar pixels in optical image, and perform joint reconstruction of the similar pixels with InSAR.

- ◇ Fusion of social media images with InSAR.

### 3-D Reconstruction from Images of Diverse Angles

Multipass InSAR is based on the concept of coherent imaging, i.e. small angle diversity between images. As the evolution of spaceborne SAR technology, abundant data are available from different viewing angles. How can we break through the limitation of coherent imaging, and perform 3-D/4-D reconstruction using images from very different angle? Two paths in this research are foreseen:

- ◇ Real SAR tomography, i.e. by back projection. This probably requires precise pixel-based matching.
- ◇ Radargrammetry with multiple images dense matching, i.e. SAR "*structure from motion*". For urban area, there have been researches on matching of images with orthogonal incidence angles (Soergel et al., 2009), and point scatterer matching (Goel and Adam, 2012b). Focus can be put on better features for dense matching. For non-urban area, there have been study on SIFT descriptor for SAR (Dellinger et al., 2015), dense matching of two images (Rodriguez Gonzalez et al., 2015), and three images matching (Raggam et al., 2010). Research is still requires on multiple images matching.

### Efficient Big Data Processing

We expect an unprecedented demand of InSAR processing in the future. Therefore, there is a necessity of developing intelligent InSAR algorithm that can select/prioritize the processing tasks. Most of the pixels in a SAR image do not undergo any significant deformation. Intelligent algorithms which can determine the hotspot of deformation in an image stack prior to the InSAR processing is very useful. The algorithms should use any easily obtainable quantities and other prior knowledge of the deformation to infer the probability of

possible large deformation in the image. One can also exploit the sparsity in spatial and temporal deformational signal, in order to reduce the number of pixels in the processing.

### **Advanced Deformation Analysis**

The current InSAR deformation estimates are still subjected to the following few points: 1) millimeter-precision of linear deformation rate over a large area requires a fairly large stack and good SNR of the scatterers; and 2) InSAR estimates are relative, although there are pioneer studies of absolute positioning and deformation, e.g. SAR imaging geodesy (Eineder et al., 2011; Cong et al., 2012; Balss et al., 2014), using active transponders (Mahapatra et al., 2014), and geodetic TomoSAR (Zhu et al., 2015c). Future research can be focused on:

- ◇ Geodetic InSAR combining absolute geodetic measurement and SAR interferometric estimates.
- ◇ Combining subsurface measurements such as seismic data with InSAR, for modelling the geophysical behaviour of volcanic region from inside and outside.
- ◇ Retrieval higher level information from the phase history, e.g. classification of different motion behaviour, and predict risk level by combining the prior knowledge of the scattering object from optical image.



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# List of Figures

- 2.1 Plane wave approximation of the interference of two SAR images. The red and blue lines are the iso-range lines with interval of half wavelength of the two images respectively. The iso-range lines intersect each other, creating an interference pattern. The background shows the interferometric phase of the pattern. The topography is mapped into interferometric phase depends on the height of the point. 5
- 2.2 Illustration of systematic shift of the multipass InSAR position estimates in UTM coordinate. The two black dots indicate the true positions of two scatterers which are on the same height w.r.t. a reference surface. The two scatterers are located at the near range and far range of the SAR image, respectively. Their incidence angles  $\theta_{near}$  and  $\theta_{far}$  are different, e.g. as many as one degree for TS-X spotlight images. Due to the unknown height of the reference point, the scatterers are shifted to the red points along their elevation directions, respectively. The scatterers are equally shifted in height, but scaled in the ground range direction due to their different elevation directions. This ground range scaling will translates into both north and east direction in the UTM coordinate. Therefore, any relative InSAR position estimate is shifted in three directions, with additional horizontal 2-D scaling. 7
- 2.3 Maximum ground range scaling (per meter of the height of reference point) as a function of the near range incidence angle and the incidence angle diversity between near range and far range. For a typical image stack of  $40^\circ$  incidence angle and  $1^\circ$  diversity, the scaling is  $3.5 \text{ cm/m}/^\circ$ . However, the scaling will become  $15 \text{ cm/m}/^\circ$  if the incidence angle becomes  $20^\circ$ , which will cause 1.5 m of scaling if the unknown height of the reference point is 10 m. 8
- 3.1 Spatial-temporal baseline plot of (a) single master InSAR stack, e.g. PSI, and (b) SBAS InSAR stack. 15
- 4.1 left: Algorithm of pre-classification of non-, single-, and double-scatterers. Non-scatterers are detected by penalized likelihood ratio test. Single and double scatterers are classified using fast non-parametric methods without TomoSAR inversion, e.g. classification using mean intensity, amplitude dispersion index, etc. And right: the flowchart of the efficient TomoSAR inversion algorithm. The most important is the layer right after the PSI processing, where pixels are pre-classified, and PSI estimates are passed to TomoSAR processing as prior knowledge. 24
- 4.2 The joint distribution densities of mean intensity and the log-likelihood to the single scatterer model for pixels with left: single scatterer, and right: double scatterer. The log-likelihood is represented by the residual sum of squares normalized to  $[0 \ 1]$ . 25
- 4.3 (a) the result of the pre-classification method, (b) the result from the SLMMER algorithm as a reference, and (c) the final classification after the higher order spectral estimation. The red pixels are detected double scatterers, the green are the single scatterers, and the blue are no scatterer. 25
- 4.4 (a) Elevation estimates of Berlin using the proposed approach, and (b) a close-up over the Berlin central station. 27

- 4.5 Fusion geometry of the proposed L-shape method. The two building façades are coloured in dark grey representing the InSAR point clouds from cross-heading orbits whose line of sight directions are indicated by the two arrows. The light grey planes denote the corresponding ground level of the two point clouds. The two pairs of red and purple dots are the façade end point position to be estimated. Due to various facts, e.g. material, decorrelation, imaging geometry, etc., these end points may not be exactly from the scatterers located on the true end position of the façades, but they are still the closest match to the exact point correspondence in two cross-heading TomoSAR point clouds. 28
- 4.6 Flowchart of the proposed algorithm. The black ellipses represent input/output data. The blue rectangles are procedures/processes. 29
- 4.7 Schematic drawing of point density estimation using directional window. The coloured points are the projection of the 3-D point cloud in horizontal plane, with the colour indicating the point density (blue: low density, red: high density). The green dashed line segment is a robust fit to the 2-D projection in the selected area within the red dashed rectangle. Points within the region bounded by the red solid rectangle are considered in the directional point density estimation, while those within the black rectangle are considered in the box-car density estimation. 30
- 4.8 Comparison of point density estimated using (a) directional window, and (b) rectangle window. The colour bar applies for both (a) and (b) with the unit  $[m^{-2}]$ . For reference, (c) is the optical image of the same area from Google. The size of directional and rectangle windows, in terms of area, are identical in this comparison. The directional window emphasizes the façade point, meanwhile keeps the point density of non-façade points unchanged compared to rectangle window. It improves the detection rate of façade points. 30
- 4.9 (a) The inlier points of one line segment of an L-shape, with the black line being the façade direction obtained from the weighted Hough transform, (b) the coordinates of the inlier points projected to the façade direction, (c) the blue curve is the estimate of (b)'s point density using a rectangular window of 5m, and the red dashed rectangle and lines demonstrate the moving line fitting method, and (d) the product of the slope and the number of inliers of the fitted line using the points in the moving rectangle window along the point density estimate shown in (c). The two red crosses are the automatically detected façade start and end position at -41.38m and 37.22m, respectively. 31
- 4.10 (a) and (b): The detected L-shapes and their end points of the two input point clouds over Berlin, and (c) the matched L-shapes and end points. The L-shapes are plotted in green, and the end points are marked as red. They are overlaid on the grey scale point density image. 32
- 4.11 The first fused TomoSAR point cloud of an entire city. The result combines an ascending and a descending point cloud of Berlin. The color indicates height on WGS84 reference surface. The unit of the colorbar is meter. 32
- 4.12 Flowchart of the semantic InSAR point cloud interpretation method. The semantic classification of the InSAR point cloud is achieved by geometric co-registering the InSAR point cloud and the optical image to a reference model. 33

- 4.13 Comparison of Google optical image (left) and TS-X high resolution spotlight image (right) of a same area. A lot of details visible in the optical image are completely hidden underneath the strong reflection of the façade scatterers in the SAR image. Images are from (Zhu and Bamler, 2012). 33
- 4.14 (a) the TomoSAR point cloud of high-rise buildings, and (b) the optical point cloud of the same area. Building façades are almost invisible in the optical point cloud, while they are prominent in the TomoSAR point cloud. 34
- 4.15 Hidden point removal operator. The blue dots are the original point cloud. And the red dots are the inverted one given a viewpoint  $O$ . 35
- 4.16 Top: optical point cloud reconstructed using standard structure from motion algorithm with colour representing the height, middle: the fused point cloud combining the optical one and two TomoSAR point clouds from ascending and descending viewing angle, and bottom: a close up view of co-registered point cloud in Berlin Potsdamer Platz, where green, red, blue represent the points from optical, ascending and descending TomoSAR point cloud, respectively. The ground points were removed for a better visualization. 37
- 4.17 (a) The first spaceborne meter-resolution 3-D TomoSAR point cloud of urban area textured with colour from optical image, (b) the corresponding TomoSAR point cloud colour-coded by height, (c) and (d) zoom in to the Berlin central station and UniversitÄtsmedizin Berlin marked in dashed red ellipses in (a), respectively. 39
- 4.18 River (red) and railway (green) classified using the BoW method. 40
- 4.19 (a) Railway points extracted from the original TomoSAR point cloud, and (b) the railway points filtered using total generalized variation. The colour shows the amplitude of seasonal motion due to the thermal expansion of the railway. 41
- 4.20 Upper: the original and the filtered estimates of amplitude of seasonal deformation in blue and red, respectively, and lower: peaks detected in the derivative of the denoised deformation function along the railway direction. 41
- 4.21 Upper: the midpoint of the detected railway joint cross-section marked in green, and lower: close up view of the railway joint. The background optical image has a ground spacing of 20cm. 42
- 4.22 Left: input noisy interferogram (upper), and the final estimated fringe (lower), and right: the input noisy interferogram (upper row) and estimated fringes (lower row) at each resolution level. The estimated fringes at each resolution level are subtracted from the noisy interferogram. The residual phase is taken as the input for the next resolution level with smaller window size. 44
- 4.23 Comparison of three covariance matrix estimators under three different observation cases; 1st row: complex circular Gaussian, 2nd row: complex t-distribution with one degree of freedom, and 3rd row: nonstationary complex t-distribution with one degree of freedom. (a) and (b) show the results of the exponential and constant coherence matrices, respectively. In both (a) and (b), 1st column: MLE (under Gaussian), 2nd column: M-estimator with t-distribution weighting, and 3rd column: rank M-estimator with t-distribution weighting. 46

- 4.24 (a) and (b): standard deviation of the coherence estimates of Gaussian MLE and the proposed robust estimator for complex t-distributed data with degree of freedom of 2.5 and 5, respectively, (c) and (d): bias of the coherence estimates. 100 looks is used in the coherence estimation. 47
- 4.25 Bias (left) and standard deviation (right) of the coherence estimates of Gaussian MLE and the proposed robust estimator as functions of degree of freedom of the complex t-distribution, for true coherence = 0.2. 100 looks is used in the coherence estimation. 48
- 4.26 Comparison of the coherence estimates using different orders of rank. The left subfigure shows the simulated interferometric phase with a constant coherence of 0.5. The other subfigures are the coherence estimates using rank 0 to 2. Rank 0 refers to using the original single look complex pixels. The colour bar indicates the coherence. Rank 0 heavily underestimate the coherence. Rank 1 greatly reduced the underestimation. Rank 2 is able to eliminate all the underestimation, but the variance of the coherence estimates also increases as the rank increases. 49
- 4.27 Comparison of the standard deviation of the linear deformation rates estimated by the robust estimators and the ordinary MLEs. The left and right subfigures correspond to the PS and DS estimators, respectively. In both subfigure, the green curves are the results of the proposed robust estimators, and the blue curves are those of the original MLEs; the dashed curves are the estimators performing on contamination-free observations, and the solid curves are the results for contaminated data. The blue dashed curves in all the subfigures are the MLEs under a complex circular Gaussian distribution; hence, they are the optimal estimates. The robust estimators improve the efficiency by a factor of 7 to 35 for contaminated data (the squared ratio of the solid curves) while still keep a relative efficiency of 70% - 80% for contamination-free data (the squared ratio of the dashed curves). 52
- 4.28 (a) the test area with significant non-linear motion is marked by the red rectangle. The red cross is the reference point. (b) the deformation phase history of the pixel marked by yellow cross in (a), where the black dots are the wrapped deformation phases and its duplicates by adding and subtracting  $2\pi$ , the black curve is a non-linear fit to the phase history, and the red and blue lines are linear fittings using RIO and MLE, respectively. 53
- 4.29 (a) linear deformation rate estimates by robust estimator and ordinary MLE, and (b) the bias of the robust and non-robust estimates to the "ground truth" (linear deformation rate estimated considering the non-linear motion model). Units: [mm/year]. 54

- 4.30 (a) and (b) comparison of the linear deformation rate of the two test sites estimated using the proposed  $\hat{\mathbf{C}}_{RME}$  and the ordinary  $\hat{\mathbf{C}}_{MLE}$ , respectively. Identical samples were used for estimating both covariance matrices. They were adaptively selected (KS test) using ten images. This is expected to introduce a non-negligible number of outliers in the selected samples because of the low detection rate from the low number of images. Results (overlaid on the SAR intensity images) are specific to a local reference point. A homogenous deformation rate is expected because of the small size of the test area. The same ordinary DS MLE was employed for the linear deformation rate estimations in both cases. And (c) and (d) the corresponding histograms of the linear deformation rates in (a) and (b), respectively. In (d), many local peaks appear almost uniformly in the search range, most of which should not correspond to the deformation signals. 55
- 4.31 Linear deformation rate of the super volcano Campi Flegrei, as retrieved using the proposed robust InSAR optimization framework that combines both PS and DS in a stack of 33 TS-X high-resolution spotlight images. Tukey's biweight function is set to be the robust loss function, and the sign covariance matrix is employed as the robust covariance matrix estimate. RIO retrieves 15 times more scatterers than those using only PS only. This number should also be valid for standard PS+DS methods, e.g. SqueeSAR. However, RIO greatly reduced the salt and pepper noise probably caused by unmodeled phase and bad covariance estimation. This can be seen in the small area comparison in which plots the results for the two areas marked by red rectangles. 56
- 4.32 Zoom in of the two areas marked by red rectangles in 4.31. 57



# List of Tables

1	Some typical loss functions, and their corresponding influence and weighting functions.	11
2	Plots of the loss functions, and their corresponding influence and weighting functions.	11
3	Overall computational cost of the proposed TomoSAR inversion method (using the test area shown in Figure 4.3 as the example), compared to using SL1MMER or SVD-Wiener only.	26
4	Comparison of RIO framework with the state-of-the-art	43
5	Algorithm of the robust estimator for single-look DS observations	51

# Acknowledgement

This work is absolutely not possible without the help of many people. First of all, my greatest gratitude goes to my supervisor Prof. Xiaoxiang Zhu, for not only but especially, sharing her comprehensive knowledge of multiple disciplines and providing unparalleled supervision. Her standard on research quality has influenced and changed many aspects of my life. Her previous excellent researches have been a perfect platform without which my work will not be visible to the community. I am grateful to her trust and support without which I could not finish this thesis after the scholarship funding period. She stresses not only scientific research but also the integrity of the team. She carefully distributes topics among the team members. I made good friends with all the team members, and feel the sense of belonging to the SiPEO team. She has also been a friend, a mentor, and a role model. As a friend, she showed great support to many aspects of my life in Germany; as a mentor, she guided me and made me a better person; and as a role model, she is a visionary and a practitioner. I am still being encouraged by her everyday passion of conducting hands-on research herself.

The same sincere gratitude goes to my supervisor Prof. Richard Bamler, for not only the same quality as Prof. Zhu but especially, his capacity of understanding and generousness. This is not only limited to scientific discussions. He does not only hear what you said, but also to what you have not said. He is a great researcher. His article "SAR Interferometry" is the sanctuary of my research in SAR signal processing, because all sorts of help and emergency assistant can be found. He has also played many other important roles. He brought laughter to us with his well-integrated science-in-life joke; he searched cheap solutions for our health insurance; and he urged me on my research from time to time with gentle force, and so on. I remember the conversation in the kitchen "none of us are tall" "but together we are strong".

Sincere gratitude also gives to Prof. Uwe Sörgel from TU Darmstadt for being the third reviewer of this thesis. His fast response to the request also plays an important role in finishing this thesis. He is a significant figure in the SAR community. His contributions in urban remote sensing have been remarkable. His scientific work together with his continuous attention in our works have encouraged me.

This thesis is also absolutely not possible without the great support from the colleagues of DLR-IMF, TUM-LMF, TUM-SiPEO, especially the SAR department led by Prof. Michael Eineder. I benefited a lot from the DLR-IMF's PSI-GENESIS, and now the IWAP system. Therefore, the whole InSAR team deserves a warm thanks, including the team leader Nico Adam, and Fernando Rodriguez, Christian Minet, Alessandro Parizzi, Dr. Ramon Brcic, Franz Ulmer, Dr. Kanika Goel, Robert Shau, and so on.

I also thank other individuals who provided support and data, including Dr. Stefan Gernhardt, Dr. Stefan Auer, Dr. Helko Breit, Gerald Baier, Dr. Michael Bäßler, Dr. Pablo d'Angelo, Dr. Ruppert Müller, Dr. Jiaojiao Tian, Dr. Jian Xu, Dr. Andrian Doicu, Dr. Xiaoying Cong, Nan Ge, Muhammad Shahzad, Claas Grohnfeldt, Dr. Heiko Hirschmüller, and so on.

Last but not least, my gratitude to my family is beyond any language. Every bit of my life right now traces back to every bit of love they offered me on my path.



# Appendices

- A Wang Y, Zhu X, Bamler R (2014) An Efficient Tomographic Inversion Approach for Urban Mapping Using Meter Resolution SAR Image Stacks. IEEE Geoscience and Remote Sensing Letters, 11 (7): 1250 - 1254.**

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# An Efficient Tomographic Inversion Approach for Urban Mapping Using Meter Resolution SAR Image Stacks

Yuanyuan Wang, Xiao Xiang Zhu, and Richard Bamler

**Abstract**—This letter describes an efficient approach of multi-dimensional synthetic aperture radar (SAR) imaging for urban mapping. The proposed approach is an integration of tomographic SAR inversion and the well-known persistent scatterer interferometry (PSI). It consists of three steps: first, a global estimation of the topography and motion parameters using efficient algorithms such as PSI; second, a single and double scatterer discrimination step based on the results of the first step; finally, a tomographic SAR inversion, which is performed on the preclassified double scatterers, using the prior knowledge obtained in the first step, retrieving the topography and motion parameters of both scatterers. The proposed approach has been tested on a dozen of TerraSAR-X high-resolution spotlight image stacks. In this letter, examples from Las Vegas and Berlin are presented. The results are comparable with the one obtained by the most computationally expensive tomographic SAR algorithms (e.g., SLIMMER) only and saves computational time by a factor of 50.

**Index Terms**—SAR tomography (TomoSAR), SLIMMER, synthetic aperture radar (SAR), TerraSAR-X, urban mapping.

## I. INTRODUCTION

### A. TomoSAR and PSI

**T**OMOGRAPHIC synthetic aperture radar (SAR) inversion includes SAR tomography (TomoSAR) [1] and differential TomoSAR (D-TomoSAR) [2]–[4]. Because of their lay-over resolving capability, they are often used for urban 3-D topographic reconstruction, forest structure imaging, and ground or building deformation monitoring. As an extension of TomoSAR, D-TomoSAR introduces an additional motion term in the TomoSAR system model, allowing the retrieval of the motion parameters of each scatterer in a resolution cell. The D-TomoSAR system model can be expressed as follows:

$$g_n = \int_{\Delta s} \gamma(s) \exp \left( -j2\pi \left( \xi_n s + \frac{2d(s, t_n)}{\lambda} \right) \right) ds \quad (1)$$

Manuscript received May 9, 2013; revised July 15, 2013 and September 3, 2013; accepted September 26, 2013. This work was supported in part by the International Graduate School of Science and Engineering of Technische Universität München under Project 6.08 4D City, by the Helmholtz Association under the framework of the Young Investigators Group “SiPEO,” and by the German Research Foundation (DFG) under Förderkennzeichen BA2033/3-1.

Y. Wang is with the Lehrstuhl für Methodik der Fernerkundung, Technische Universität München, 80333 Munich, Germany (e-mail: wang@bv.tum.de).

X. X. Zhu and R. Bamler are with the Lehrstuhl für Methodik der Fernerkundung, Technische Universität München, 80333 Munich, Germany, and also with the Remote Sensing Technology Institute (IMF), German Aerospace Center (DLR), 82234 Weßling, Germany (e-mail: xiao.zhu@dlr.de).

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Digital Object Identifier 10.1109/LGRS.2013.2290833

where  $g_n$  is the complex pixel value in the  $n$ th single-look complex SAR image,  $s$  represents the elevation direction,  $d(s, t_n)$  is the displacement as a function of the elevation and the acquisition time  $t_n$ ,  $\lambda$  is the wavelength of the SAR system,  $\xi_n = -2b_n/(\lambda r)$ , which is the so-called “spatial frequency” and is proportional to the baseline  $b_n$  and range  $r$ , and  $\gamma(s)$  is the complex reflectivity profile along the elevation direction of that pixel. Solving the reflectivity  $\gamma(s)$  is essentially a spectral estimation problem. In case of solving it parametrically, the solution is, in addition, subject to a multidimensional search in the elevation and motion parameters solution space.

Persistent scatterer interferometry (PSI) [5] assumes a maximum single dominant scatterer in a pixel. Under such assumption,  $\gamma(s)$  in (1) is modeled as a delta function with a constant amplitude. Assuming linear and seasonal motion models (common in urban area), the PSI system model can be simplified as

$$g_n = A \exp(-j2\pi(\xi_n s + \eta_n v + \omega_n a)) \quad (2)$$

where  $A$  is the amplitude of the point scatterer, which is usually dropped in the estimation,  $\eta_n = 2t_n/\lambda$  is the temporal frequency modeling linear movement, and  $\omega_n = 2\sin(2\pi(t_n - t_0))/\lambda$  is the temporal frequency modeling seasonal motion.  $v$  and  $a$  are the velocity of linear motion and the amplitude of periodic motion, respectively. Solving the unknown elevation and the motion parameters is to match the phase on both sides of (2), which can be efficiently solved by maximizing the ensemble coherence (periodogram) [6]. Because of the simplification in the system model, PSI is an efficient method for deformation monitoring of large areas.

### B. Integration of TomoSAR and PSI

As the resolution of SAR images goes higher, it provides us enormous details on one hand, and on the other hand, it increases the computational burden. This is particularly true for monitoring an entire urban area using D-TomoSAR because it is a spectral estimation problem whose computational efficiency is mainly restricted by the following: 1) dimension of the spectral estimation problem for each pixel. Usually, 3-D is applied (elevation, linear motion, and periodic motion) [7]; 2) extent of the parameter solution range, i.e., the search range in each dimension of the solution space; and 3) the spectral estimation algorithms being applied, such as periodogram (economic), SVD-Wiener [4] (a little bit more expensive) and SLIMMER [8] (much more expensive).

To reduce the computation, we propose to integrate PSI into the D-TomoSAR inversion, with these considerations. 1) PSI

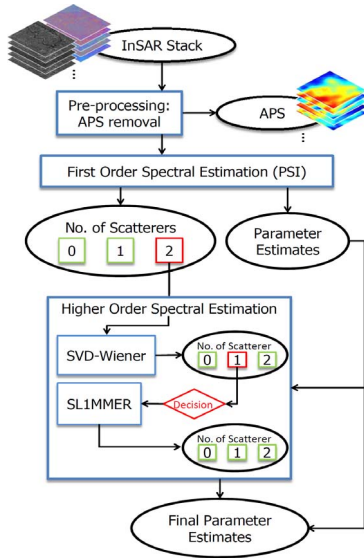


Fig. 1. Processing flowchart. Black ellipses represent input/output data. Blue rectangles represent processes/algorithms. The red diamond represents decision. The green rectangles in the black ellipses represent accepted results, whereas those red ones represent results that need to be passed to the next step.

needs to provide a reasonable atmospheric phase screen (APS) estimation; 2) PSI should provide the prior knowledge about the dimension of the problem (i.e., the motion model) and the search range in each dimension of the solution space; 3) More expensive methods such as SVD-Wiener and SLIMMER shall only be applied to the pixels that probably contain more than one scatterer; and 4) Algorithms with superresolving capability such as SLIMMER shall only be applied when superresolution is required.

### C. Summary

The proposed TomoSAR inversion approach is an integration of PSI and D-TomoSAR. The processing flowchart is shown in Fig. 1. It follows three steps: first-order spectral estimation, single and double scatterer(s) discrimination, and higher order spectral estimation. A near-operational processor is implemented by combining the PSI, SVD-Wiener, and SLIMMER developed in the Remote Sensing Technology Institute of the German Aerospace Center (DLR).

The rest of the letter will describe each step in this approach. The details of PSI, SVD-Wiener, and SLIMMER will not be addressed and can be found in [4], [5], and [8]. Focus will be put on the integration of PSI and D-TomoSAR, particularly the discrimination of pixels with single or double scatterer(s). Results using TerraSAR-X high-resolution spotlight data are presented in Section III.

## II. TOMOSAR PROCESSING CHAIN

The proposed approach is summarized in Fig. 1. First, APS is removed from the coregistered image stack in the PSI-type preprocessing; then, first-order spectral estimation (i.e., single-scatterer model) is applied on all the pixels, obtaining the estimates of the model parameters. Based on these estimates, amplitude dispersion index (ADI), and pixel brightness, a pre-discrimination of the number of scatterers in each pixel is then obtained. As we limit the maximum number of scatterers to two, all the pixels shall fall into one of the three classes: no,

single, or double scatterer(s). The prediscrimination follows two steps: The Bayesian decision rule is first applied to discard pixels with no scatterer and followed by a support vector machine (SVM) to separate pixels with single scatterer and double scatterers. The pixels detected with double scatterers (upper red box in Fig. 1) are passed to higher order spectral estimation. They are first processed with SVD-Wiener. Some of them are confirmed as pixels containing double scatterers, and the rest are rejected as having either single or no scatterer. Because SVD-Wiener has almost no superresolution capability, among those pixels rejected as single scatterers (lower red box in Fig. 1), some actually contain double scatterers. Therefore, after a decision (red diamond in Fig. 1) on the elevation distance, some of them are passed to the SLIMMER algorithm with superresolution capability. The following text in this section will explain each step in detail.

### A. Preprocessing

The purpose of preprocessing is to estimate and further remove the APS from a coregistered complex SAR image stack. The image stacking is done by DLR's PSI-GENESIS system [9]. The APS is removed using a similar approach as that in standard PSI processing and the "spatial difference" method mentioned in [10]. Since the APS is spatially smooth varying, preprocessing is performed on the downsampled image stacks to reduce computational cost. Note that the pixel selection and parameter estimates in the preprocessing do not affect/replace any procedures in the following D-TomoSAR inversion steps. The basic preprocessing procedures are listed as follows.

- 1) Downsample images in azimuth and range direction.
- 2) Build redundant network from spatially differential measurement in the downsampled image stacks.
- 3) Estimate model parameters of the differential measurements in the network, assuming a single-scatterer model with linear and seasonal motion.
- 4) Integrate differential topography estimates and remove the corresponding phase contribution from the original interferometric phase.
- 5) Unwrap the remaining residual phase in azimuth and range direction.
- 6) Apply a temporal high-pass filter and a spatial low-pass filter to the unwrapped phase stack. This results to the APS in the downsampled image size.
- 7) Upsample the APS to the original image size and remove from the original phase.

### B. First-Order Spectral Estimation

The first-order spectral estimation refers to a special case of the D-TomoSAR problem considering only a single scatterer in each range-azimuth pixel. In the proposed processing chain, it is solved by maximizing the periodogram [11] of the amplitude-normalized measurements [5], i.e.,

$$\{\hat{s}, \hat{v}, \hat{a}\} = \arg \max \left\{ \left| \frac{1}{N} \sum_{n=1}^N e^{-j2\pi(\xi_n s + \eta_n v + \omega_n a)} \frac{g_n^*}{|g_n|} \right| \right\} \quad (3)$$

where  $N$  is the total number of interferograms, and  $*$  stands for the complex conjugate operator.

After applying the estimation on every pixel, we obtain the estimates of the elevation and motion parameters related to a reference point that we assume stable. Median and maximum filters are applied on the range–azimuth image of the estimates to obtain the prior for higher order spectral estimation. In addition, the following quantities are also recorded: 1) pixel brightness; 2) ADI; and 3) likelihoods of the single- and no-scatterer models w.r.t. the measurements  $g_n$ .

Under a circular Gaussian assumption of the measurements with covariance  $\varepsilon$ , the log likelihood of the single-scatterer model, i.e.,  $H_1$ , is simply the sum of squared residuals [4], i.e.,

$$\ln p(\mathbf{g}|H_1) = \ln \left( \frac{1}{\pi^N |\varepsilon|^{2N}} \right) - \frac{1}{|\varepsilon|^2} \sum_{i=1}^N \left| g_n - \hat{A} \exp(-j2\pi(\xi_n \hat{s} + \eta_n \hat{v} + \omega_n \hat{a})) \right|^2 \quad (4)$$

and for the no-scatterer model, i.e.,  $H_0$ , it is the sum of squared measurements, i.e.,

$$\ln p(\mathbf{g}|H_0) = \ln \left( \frac{1}{\pi^N |\varepsilon|^{2N}} \right) - \frac{1}{|\varepsilon|^2} \sum_{i=1}^N |g_n|^2. \quad (5)$$

### C. Number of Scatterers Discrimination

Discriminating pixels with different numbers of scatterers is a classification problem. One of the optimum detection strategies is the likelihood ratio test, if the assumption of the underlying models is correct. However, such detection strategy requires the likelihood to the double-scatterer model, i.e.,  $H_2$ , and hence is not feasible before the TomoSAR inversion. Therefore, in order to prediscriminate the pixels before a TomoSAR inversion, our method is a two-step approach, comprising both theoretical and empirical parts.

First, pixels without scatterer are discriminated following the Bayesian decision rule with penalized likelihood criterion, i.e.,

$$\frac{p(\mathbf{g}|H_0)}{p(\mathbf{g}|H_1) \exp(-\text{AIC})} \frac{P_0}{P_{1,2}} \stackrel{H_0}{>} 1. \quad (6)$$

The  $H_0$  and  $H_1$  likelihoods of each pixel are compared, taking into account the Akaike information criterion (AIC) [4].  $P_0$  and  $P_{1,2}$  are the prior probabilities of pixels with no scatterer and with single or double scatterer(s), respectively. Close to 100% detection rate and 0% false alarm rate can be achieved in this stage because the  $H_1$  likelihoods of the pixels with double scatterers are much higher than their  $H_0$  likelihoods. This is proven in the real data experiment presented in Section III-A.

The remaining pixels need to be further classified into single and double scatterer(s). Since the  $H_2$  likelihood is not available at this stage, a few other easily accessible features associated with each pixel are used for the classification. They are ADI,  $H_1$  likelihood, and pixel mean intensity, with these three reasons.

- 1) Pixels with single scatterer experience lower ADI since the amplitudes of pixels subjected to only a single dominant scatterer do not show pronounced fluctuations as a function of baseline.
- 2) The  $H_1$  likelihood tends to be lower if multiple scatterers are present.
- 3) The aforementioned two aspects are based on a reasonable signal-to-noise ratio range. Therefore, only relatively

bright pixels should be selected as candidates of pixels with double scatterers.

Without the analytical model of the joint probability density function of the aforementioned three quantities, we employ the SVM to find a separation of these two classes. Considering our situation of missing classification ground truth, the usage of the SVM basically follows these points.

- The training data of the SVM classifier are the classification results obtained by SL1MMER (theoretically, the closest to ground truth) [12] on a small and representative test area. The SVM classifier takes into account three features: ADI,  $H_1$  likelihood, and pixel mean intensity.
- A nonlinear SVM classifier is applied. A Gaussian kernel is used, which is suggested in [13] as a good first try.
- The optimum parameter setting is determined by the following.
  - 1) Train the classifier using 50% of the pixels in the small test area.
  - 2) Classify all the pixels in the test area using the trained classifier.
  - 3) Repeat this with a geometric sequence of the parameters and find the parameters that give the highest detection rate at a constant false alarm rate.
- The trained SVM classifier is then used to classify all the pixels in the whole scene, based on the three features of each pixel.

### D. Higher Order Spectral Estimation

(SVD-Wiener, SL1MMER)

All pixels classified as double scatterers are processed using SVD-Wiener first, taking into account the prior knowledge of the estimates obtained in the first-order spectral estimation. In the SVD-Wiener processing, these pixels are tested through model order selection and are classified as no, single, or double scatterer(s) again. Two layers of elevation and motion parameter estimates are retrieved from the pixels confirmed with double scatterers. The lower layer of elevation estimates is interpolated in range–azimuth dimension to get a ground surface height  $s_0$  for each pixel (the index of the pixel is ignored in  $s_0$ ).

Since SVD-Wiener has almost no superresolution capability, double scatterers with elevation distance shorter than the Rayleigh resolution  $\rho_s = \lambda r / 2\Delta b$  [1] ( $r$ : slant range;  $\Delta b$ : elevation aperture size) will be misclassified as single scatterer. Therefore, among the detected pixels with single scatterer, those whose elevation estimates with respect to its corresponding ground surface height are smaller than  $\rho_s$  (in practice, we allow some buffer and set it to be  $1.5\rho_s$ ) are potential pixels that would require superresolution. In this case, SL1MMER is applied. This decision is marked as the red diamond in the processing flowchart in Fig. 1. In the SL1MMER processing, the search range is restricted to  $[s_0 - s_0 + 1.5\rho_s]$ .

## III. VALIDATION WITH REAL DATA

### A. Experiments on Number of Scatterers Discrimination

This section will evaluate the performance of the scatterer discriminator over a test area in downtown Las Vegas shown in Fig. 2. This stack contains 25 TerraSAR-X high-resolution spotlight images. Since the missing of ground truth, the SL1MMER result is taken as reference for comparison.



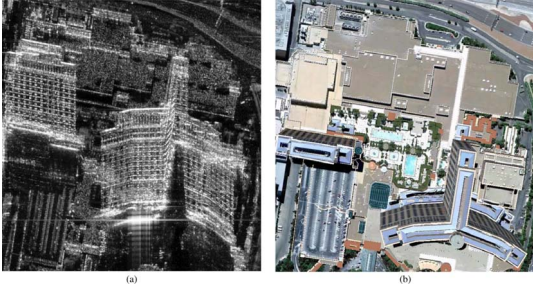


Fig. 2. (a) Amplitude image of the test area in downtown Las Vegas. (b) Optical image from Google Earth of the same area.

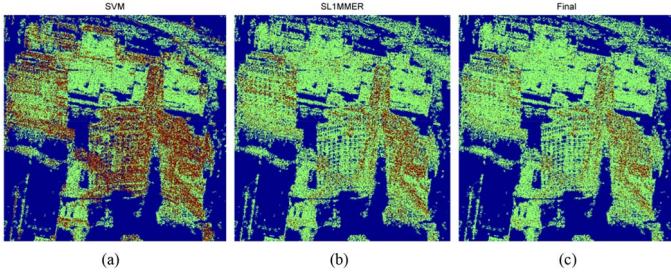


Fig. 3. (a) Preclassification result of the proposed method (likelihood ratio test + SVM). (b) Result from the SLIMMER algorithm. (c) Final classification after the higher order spectral estimation. Red represents pixels with double scatterers, green represents pixels with single scatterer, and blue represents pixels without coherent scatterer.

First, the pixels without scatterer are determined according to (6). The prior probabilities  $P_0$  and  $P_{1,2}$  are set to be both 50% in the experiment. Then, the remaining pixels are separated using an SVM into containing either single or double scatterer(s). Fig. 3(a) and (b) shows a comparison of the SVM and SLIMMER classification results, with red representing pixels with double scatterers, green representing pixels with single scatterer, and blue representing pixels with no scatterer.

In Fig. 3(a), 47% of the pixels are detected as no scatterers, 36% are detected as single scatterers, and the remaining 17% are detected as double scatterers. The classification of pixels with no scatterer is almost 100% correct compared with the SLIMMER result. In the rest of the pixels, the detection rate of pixels with double scatterers is 72.4%, with a false alarm rate of 25.7%, which is why much more pixels are detected as double scatterers. However, higher detection rate (consequently, higher false alarm rate) is still favored in the processing because the falsely detected pixels with double scatterers can still be corrected in the higher order spectral estimation, but not for the misdetected pixels with double scatterers. In order to fully characterize the proposed single and double scatterer(s) detector, its receiver operating characteristic (ROC) curve is plotted in Fig. 4. The blue dots are the experimental results, and the green curve is an analytical fit. Depending on the processing power, the threshold can be shifted toward higher detection rate, affording higher false alarm rate, and, therefore, more processing effort by more frequently applying SVD-Wiener and SLIMMER.

Next, SVD-Wiener is applied on the red pixels shown in Fig. 3(a). Most of them are confirmed as double scatterers, and others are rejected as no or single scatterer. Among those rejected as single scatterers, those having small elevation estimates (60 m in our case = 1.5 times of  $\rho_s$ ) w.r.t the ground surface height are handed over to SLIMMER for reprocessing.

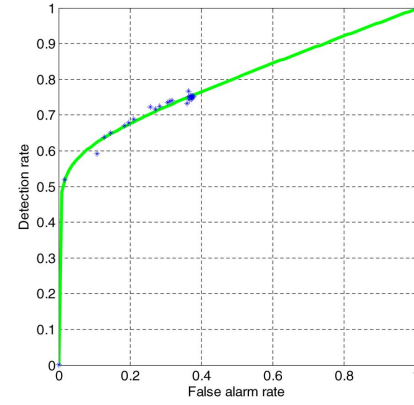


Fig. 4. ROC curve of the proposed SVM double scatterers detector. The blue dots are the best detection rates achieved in the experiments at a specific false alarm rate. The green curve is an analytical fit to the blue dots.

They account for 9% of the total pixel number in the test area. Then, the final corrected classification map of the test area can be obtained, which is shown in Fig. 3(c).

For a quantitative measure of the final classification, 98% of the pixels are correctly classified (the SLIMMER results in Fig. 3(b) as reference). The major classification error comes from the misdetection of pixels with double scatterers. This can be improved by allowing higher detection rates in the double scatterer detection. However, computation effort will also increase correspondingly since more pixels will be processed by higher order spectral estimation.

## B. Analysis of the Computational Complexity

This section will discuss the computational complexity of the proposed approach. We define  $O(1)$  to be the computation time for one multiplication. The discussion will be focused on the simple case without any motion model. To extend to a single- or a multicomponent motion model, the computational complexity exponentially increases with the number of dimensions in the system model of the problem.

In the first-order spectral estimation, the periodogram consists of multiplication of the measurement vector with the modeled phase and repeats at different elevation positions. Therefore, the computational complexity is  $O(NL)$ , where  $N$  is the number of images (usually 20–50), and  $L$  is the discretization level in the elevation direction (usually 50–200 for TerraSAR-X). In the higher order spectral estimation, SVD-Wiener needs at least  $O(N^2 + NL)$  according to [4, eq. (11)]. The SLIMMER algorithm requires a  $L_1-L_2$  norm optimization, which is extremely computationally costly, e.g., the well-known Basic Gradient Projection for Sparse Reconstruction solver [14] requires at least  $O(KML_s^2)$ , where  $K$  is the number of iterations (approximately 20),  $M$  is the number of multiplication of a specific matrix in each iteration (modest, usually several), and  $L_s$  is the elevation discretization level required for sparse reconstruction, which is about ten times greater than  $L$ . Our experience is that the SLIMMER algorithm is several hundred times slower than SVD-Wiener.

When prior knowledge of the elevation span is available,  $L$  can be restricted to a specific elevation range and hence reduce the computational cost. For instance, assuming a uniform distribution of building heights prior over the processing area, the total processing cost can be reduced to half for SVD-Wiener

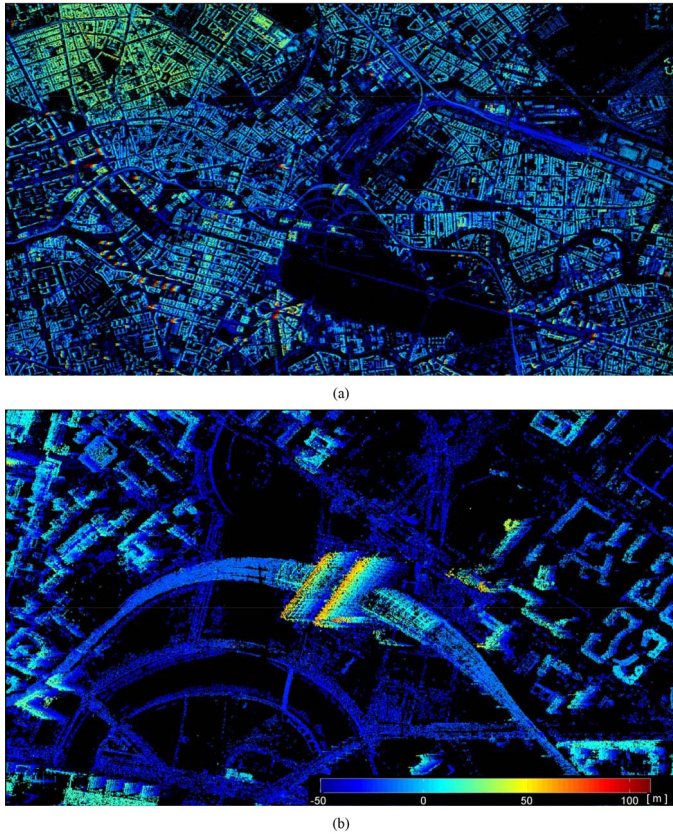


Fig. 5. (a) Elevation estimates of Berlin using the proposed approach. (b) Close-up view over the Berlin central station.

when  $L \gg N$  and to one third for SLIMMER. Therefore, it is always more economic to obtain the prior through some low-cost methods before the higher order spectral estimation.

The overall computational cost also depends on the percentage of pixels allocated to each algorithm. As an example, for the test area discussed in Section III-A, 100% of the pixels are first processed by a periodogram, 17% are then passed to SVD-Wiener, and only 9% are finally processed by SLIMMER. This small percentage together with the restriction of the search space leads to an overall speedup of about 50 compared with SLIMMER-only processing. The computational cost of the pixel preclassification using the SVM is negligible compared with the PSI plus TomoSAR processing and, hence, ignored in this analysis. Quicker processing can be achieved if no superresolution is required.

### C. Large-Area Processing: Berlin

The proposed processing chain is applied on several stacks of TerraSAR-X high-resolution spotlight images over Berlin. Fig. 5 shows the elevation estimates and a close-up view of one of the stacks, which is composed of 94 images. In the close-up view of the Berlin central station, the proposed processing chain is able to catch most of the structure.

## IV. CONCLUSION AND OUTLOOK

This letter has proposed an efficient TomoSAR inversion approach for meter resolution SAR image stacks of urban areas. It is the integration of three spectral estimation algorithms: periodogram, SVD-Wiener, and SLIMMER. It has been tested

on a number of image stacks of urban areas. Reliable results are obtained, and a speedup factor of over 50 is confirmed. The computational cost is saved in two aspects: prior knowledge of the solution search range is obtained through low-cost periodogram, and only pixels with more than one strong scatterer are processed using SVD-Wiener, and SLIMMER is only applied to pixels that require superresolution. The resulting very high resolution (VHR) spaceborne TomoSAR point clouds can be used for multiple applications. For example, individual building monitoring over the whole city, fusion of point cloud from different angles [15], and city 3-D model reconstruction from the point clouds [16].

## ACKNOWLEDGMENT

The authors thank the Gauss Centre for Supercomputing (GCS) e.V. for funding this project by providing computing time on the GCS Supercomputer SuperMUC at the Leibniz Supercomputing Centre (LRZ).

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# Appendices

- B Wang Y, Zhu X (2015a) Automatic Feature-based Geometric Fusion of Multi-view TomoSAR Point Clouds in Urban Area. IEEE Journal of Selected Topics in Applied Earth Observation and Remote Sensing, 8 (3): 953 - 965.**

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# Automatic Feature-Based Geometric Fusion of Multiview TomoSAR Point Clouds in Urban Area

Yuanyuan Wang, *Student Member, IEEE*, and Xiao Xiang Zhu, *Senior Member, IEEE*

**Abstract**—Interferometric synthetic aperture radar (InSAR) techniques, such as persistent scatterer interferometry (PSI) or SAR tomography (TomoSAR), deliver three-dimensional (3-D) point clouds of the scatterers' positions together with their motion information relative to a reference point. Due to the SAR side-looking geometry, minimum of two point clouds from cross-heading orbits, i.e., ascending and descending, are required to achieve a complete monitoring over an urban area. However, these two point clouds are usually not coregistered due to their different reference points with unknown 3-D positions. In general, no exact identical points from the same physical object can be found in such two point clouds. This article describes a robust algorithm for fusing such two point clouds of urban areas. The contribution of this paper is finding the theoretically exact point correspondence, which is the end positions of façades, where the two point clouds close. We explicitly define this algorithm as “L-shape detection and matching,” in this paper, because the façades commonly appear as L-shapes in InSAR point cloud. This algorithm introduces a few important features for a reliable result, including point density estimation using adaptive directional window for better façade points detection and L-shape extraction using weighed Hough transform. The algorithm is fully automatic. Its accuracy is evaluated using simulated data. Furthermore, the proposed method is applied on two TomoSAR point clouds over Berlin with ascending and descending geometry. The result is compared with the first PSI point cloud fusion method (S. Gernhardt and R. Bamler, “Deformation monitoring of single buildings using meter-resolution SAR data in PSI,” *ISPRS J. Photogramm. Remote Sens.*, vol. 73, pp. 68–79, 2012.) for urban area. Submeter consistency is achieved.

**Index Terms**—Point cloud fusion, SAR tomography (TomoSAR), synthetic aperture radar (SAR), TerraSAR-X.

Manuscript received May 30, 2014; revised August 22, 2014; accepted September 18, 2014. This work was supported in part by the International Graduate School of Science and Engineering, Technische Universität München (Project 6.08: “4D City”), in part by the Helmholtz Association under the framework of the Young Investigators Group “SiPEO” (VH-NG-1018, www.sipeco.bgu.tum.de), in part by the German Research Foundation (DFG, Förderkennzeichen BA2033/3-1), and in part by the Gauss Centre for Supercomputing e.V. (www.gauss-centre.eu) by providing computing time on the GCS Supercomputer SuperMUC at Leibniz Supercomputing Centre (LRZ, www.lrz.de).

Y. Wang is with the Helmholtz Young Investigators Group “SiPEO”, Technische Universität München, Munich 80333, Germany (e-mail: wang@bv.tum.de).

X. Zhu is with the Helmholtz Young Investigators Group “SiPEO”, Technische Universität München, Munich 80333, Germany, and also with the Remote Sensing Technology Institute (IMF), German Aerospace Center (DLR), Weßling 82234, Germany (e-mail: xiao.zhu@dlr.de).

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Digital Object Identifier 10.1109/JSTARS.2014.2361430

## I. INTRODUCTION

### A. TomoSAR

**T**OMOGRAPHIC SAR inversion (TomoSAR) [2]–[6] is an interferometric synthetic aperture radar (InSAR) technique that tracks down the long-term or partially coherent pixels in a stack of coregistered SAR images, which is similar to other InSAR techniques such as persistent scatterer interferometry (PSI) [7]–[10], small baseline subset (SBAS) [11]–[13], SqueeSAR [14], [15], and CEASAR [16]. They all aim at retrieving the three-dimensional (3-D) position and the parameters of the undergoing motion of point and/or volumetric scatterers.

However, fundamentally different, TomoSAR exploits possible multiple scatterers in the third dimension *elevation* of a rang-azimuth pixel—a phenomenon called *layover* in radar jargon that happens when two or more scatterers are at the same range (distance) to the sensor. For example, the closely packed vertical structures in urban areas cause severe layover. TomoSAR retrieves the elevations and motion parameters of multiple scatterers by means of spectral estimation. This is not possible with the above-mentioned single scatterer model-based methods, e.g., PSI. Therefore, TomoSAR is so far the most competent InSAR method for long-term urban monitoring [17].

Therefore, the 3-D TomoSAR point clouds derived from meter-resolution SAR image stacks are the basis of the work of this paper. The point clouds we used are obtained by *Tomo-GENESIS* [18]—a TomoSAR software of the German Aerospace Center (DLR) for large urban areas monitoring. It is developed based on the work of [5], [6], [17], [19]–[21]. It delivers, for the first time, a 5-D point cloud—3-D position plus linear deformation rate and amplitude of seasonal motion—from a stack of SAR images. The point density is up to  $10^6/\text{km}^2$ , when using TerraSAR-X high-resolution spotlight data. Such density is comparable with some light detection and ranging (LiDAR) product and enables the retrieval of the features of individual building.

### B. Fusion Geometry of Cross-Heading InSAR Point Clouds

Cross-heading orbits refer to satellite's ascending and descending flying trajectories. Due to the side-looking geometry of SAR, minimum of two TomoSAR point clouds from cross-heading orbits are required to achieve a complete monitoring over the whole area. As demonstrated in Fig. 1, point cloud derived from SAR image stacks of ascending or descending orbits provides either the front or the rear side of the same building. Such two point clouds have basically no *exact point*

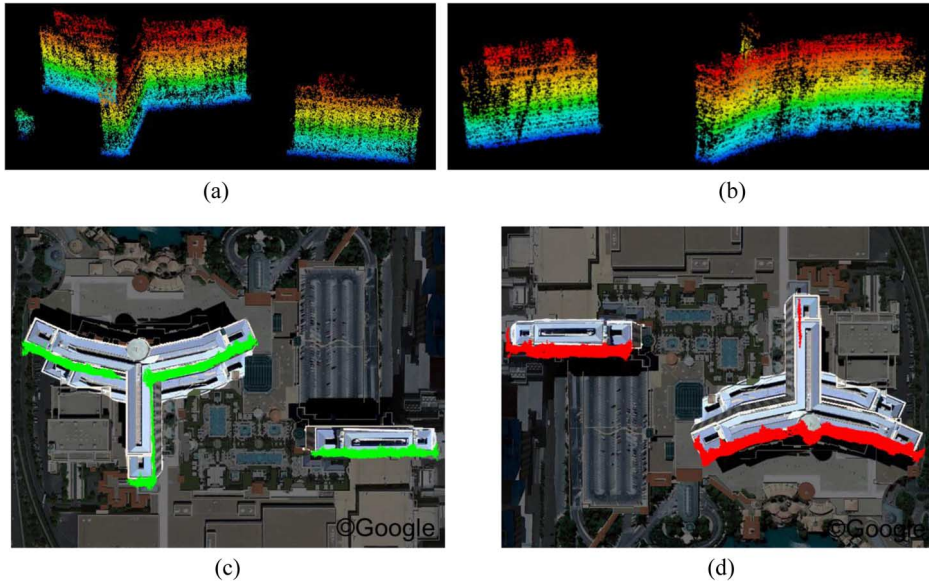


Fig. 1. TomoSAR point cloud derived from (a) an ascending image stack in Las Vegas and (b) a descending image stack of the same area. Color represents the height. (c) and (d) are the corresponding footprint of (a) and (b) on Google Map satellite images.

*correspondence*, which is referred to as the identical geolocation of the sources of two corresponding points (in the two point clouds, respectively). Examples are man-made objects like cylindrical reflector or 3-D corner reflector that could reflect radar signal in all directions. Not difficult to imagine, fusion of such point clouds is as tricky as coregistering two optical images of front and rear side of an opaque object.

These two point clouds are always relative to two different reference points with unknown heights. The uncompensated height of the reference point causes the point cloud to shift along the elevation direction. Therefore, the fusion task is to find the unknown heights of the two reference points. A schematic drawing of fusion in Universal Transverse Mercator (UTM) coordinate is illustrated in Fig. 2 (modified from [1]), where the left and right satellites indicate the ascending and descending master orbits at a fixed *east-up* plane, respectively. Perpendicular toward the reader's screen/paper is the *north* direction. The flying direction of the two satellites is indicated by the cross (fly away from reader) and dot (coming toward reader) on the satellites. They are not parallel to the north direction. The angle between the flying direction and the meridian (north) is the so-called heading angle.  $R^a$  and  $R^b$  are the range distances of the two orbits. The two black dots are the corresponding scatterers assumed to have identical position, but geocoded into different positions  $\mathbf{P}_{xyz}^a$  and  $\mathbf{P}_{xyz}^b$  due to their different and unknown reference height  $\Delta z^a$  and  $\Delta z^b$ . This also causes horizontal shifts  $\Delta xy^a$  and  $\Delta xy^b$ , which are functions of the incidence angles  $\theta^a$  and  $\theta^b$ , i.e., the horizontal shift is not constant throughout the whole point cloud. To achieve a fusion, the two scatterers have to be shifted along their elevation directions by  $\Delta S^a = \Delta z^a / \sin \theta^a$  and  $\Delta S^b = \Delta z^b / \sin \theta^b$ , respectively. The fusion model in UTM coordinate system is then [1]

$$\mathbf{P}_{xyz}^a + \frac{\Delta z^a}{\sin \theta^a} \mathbf{s}^a = \mathbf{P}_{xyz}^b + \frac{\Delta z^b}{\sin \theta^b} \mathbf{s}^b \quad (1)$$

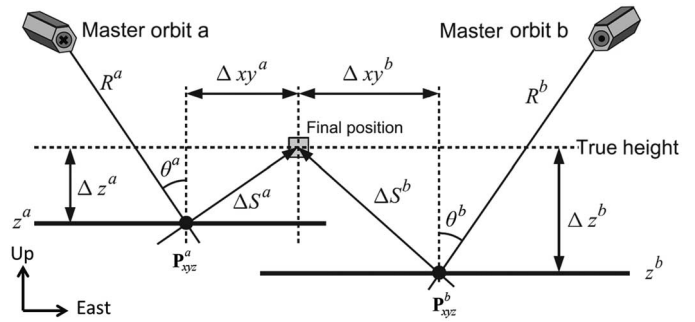


Fig. 2. Schematic representation (modified figure from [1]) of the InSAR point cloud shift in UTM coordinate due to different reference points. The reader's screen/paper is the east-up plane, and perpendicular to it away from the reader is the north direction. The two satellites indicate the ascending and descending master orbits positions. Their flying directions are indicated by the cross (fly away from reader) and dot (coming toward reader) on the satellites. They are not parallel to the north direction. The angle between the flying direction and the meridian (north) is the so-called heading angle. The two black dots are the two scatterers assumed to be identical in UTM coordinate system, but shifted in the two point clouds due to the heights  $\Delta z^a$  and  $\Delta z^b$  of their reference points.

where bold letters indicate vectors, and  $\mathbf{s}^a$  and  $\mathbf{s}^b$  are the unit vectors of the elevation directions of the two scatterers, which depends on the incidence angle  $\theta$ , and heading angle  $t$

$$\mathbf{s} = \begin{bmatrix} \cos t \cos \theta \\ -\sin t \cos \theta \\ \sin \theta \end{bmatrix}. \quad (2)$$

In another words, three equations can be written for each pair of corresponding points ( $\mathbf{P}_{xyz}^a, \mathbf{P}_{xyz}^b$ )

$$\begin{aligned} x^a - x^b + \Delta z^a \cos t^a \cot \theta^a - \Delta z^b \cos t^b \cot \theta^b &= 0 \\ y^a - y^b - \Delta z^a \sin t^a \cot \theta^a + \Delta z^b \sin t^b \cot \theta^b &= 0 \\ z^a - z^b + \Delta z^a - \Delta z^b &= 0 \end{aligned} \quad (3)$$



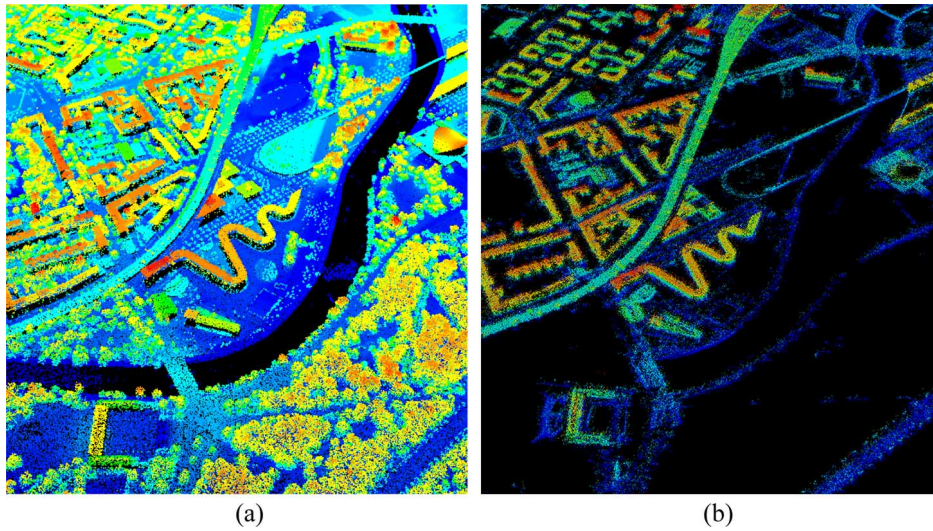


Fig. 3. Comparison of (a) LiDAR point cloud [22] and (b) TomoSAR point cloud of the same area in downtown Berlin. The color represents the height. LiDAR point cloud has, in general, better quality in terms of 3-D positioning accuracy.

where  $\Delta z^a$  and  $\Delta z^b$  are the only unknowns. However, the incidence and heading angles  $\theta$  and  $t$  are direct functions of the point current position. As described in [1], this problem is the so-called *Gauß-Helmert* model. It needs to be solved iteratively, where  $\mathbf{P}_{xyz}^{\{a,b\}}$ ,  $\theta^{\{a,b\}}$ , and  $t^{\{a,b\}}$  are updated at each iteration. The estimation is robust with many pairs of points available.

### C. Related Work and Motivation

In computer vision, the 3-D rigid point cloud coregistration methods are very well studied. For a successful coregistration, overlapping of the two point clouds is required, so that one can find common points or features that describe a common location in the point clouds. The basic approach of such task follows the procedures of feature detection, description, and matching, where so do the researches focus. The feature detection and description are usually combined. The most well-known ones are:

- 1) gradient-based features, e.g., scale invariant feature transform (SIFT) [23] and speeded up robust features (SURF) [24];
- 2) histogram-based features, e.g., histogram of gradients (HoG) [25] and point feature histogram (PFH) [26]; and
- 3) surface normal-based features, e.g., spin-image [27] and normal-aligned radial feature (NARF) [28].

Here, we just name a few. Many of their variants are also highly used. The matching algorithms, according to the optimization methods used, can be classified as either global or local. As pointed out in [26], the most famous algorithms of the first type are generic and evolutionary algorithms [29], [30]. Not surprisingly, most of the study is the second type, due to the computational complexity of the first type. The most well-known one is the iterative closest point (ICP) [31], [32]. Other highly cited works are mostly variants or improvements in the original ICP. For instance, to find better point correspondence using feature descriptors [26], [33], [34], to find better initial

transformation [34], [35], and to use nonlinear optimization methods in ICP [36], [37].

In the remote sensing community, these techniques are employed for coregistration of LiDAR or optical image derived point clouds [38]–[42]. However, no attempt has been made for fusion of PSI or TomoSAR-derived point clouds of urban areas. This is especially true for two point clouds generated from data stacks of cross-heading orbits, due to the lack of exact point correspondence. Another reason prevents the above-mentioned methods from being directly applicable is the relatively worse accuracy of InSAR point clouds compared with LiDAR or optical image-derived ones. An example is made in Fig. 3, which is the comparison of LiDAR and TomoSAR point cloud of the same area in Berlin.

The first attempt of such task is described in [1] using PSI point clouds. It is essentially a 3-D surfaces matching that relays on removing the façade points, where the point correspondence is most unlikely to appear. The point correspondences are found by searching closely space point pairs on surface. Its accuracy is limited, since the exact point correspondence is not addressed. The computational efficiency is compromised when applying to TomoSAR point clouds [43]. Practically, only 10% of the non-façade points are used, since a TomoSAR point cloud is usually much denser than PSIs’.

Therefore, the initiative of the proposed method is to make use of the rich façade information in TomoSAR point clouds in order to derive the theoretically exact point correspondence and to develop a computationally efficient algorithm that can handle large point clouds. Finally, we can obtain a 3-D point cloud associated with the movement information covering an entire city.

As no exact point correspondence can be found among the points of the two point clouds, we will have to look for higher level features. Studies such as [44]–[46] have already suggested that quadrilateral buildings appear as “L-shapes” in SAR images. As a result, the end points of the L-shape, i.e., the building edges where the two point clouds close, can be used as the



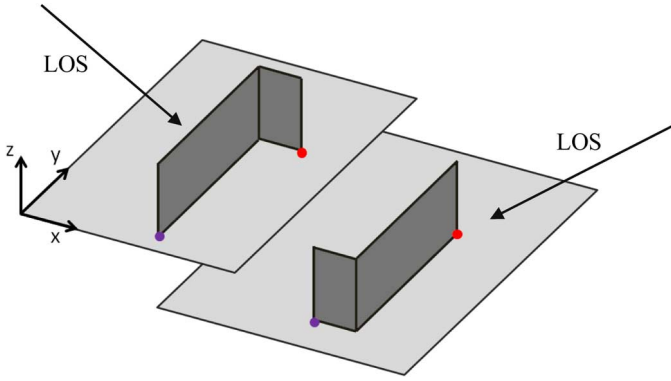


Fig. 4. Fusion geometry of the proposed L-shape method. The two building façades are colored in dark gray representing the geocoded point clouds from cross-heading orbits whose LOS directions are indicated by the two arrows. The light gray planes denote the corresponding ground level of the two point clouds. The two pairs red and purple dots are the façade end point position to be estimated.

exact point correspondence. Described in Fig. 4, the L-shape end points are indicated by the two pairs of red and purple dots. The two building façades colored in dark gray represent the point clouds from ascending and descending orbits whose line of sight (LOS) directions are indicated by the two arrows, respectively. The light gray planes denote the corresponding ground levels of the two point clouds.

The coregistration is then broken down to the task of detecting the L-shapes' end points, estimating their 3-D positions as the descriptor, and finally matching the two reduced point clouds. To achieve this, some robust procedures are designed to extract façade points, estimate the positions of the façade edges, and then find correspondence in the two point clouds.

## II. FEATURE-BASED INSAR POINT CLOUD FUSION

The flowchart of the algorithm is plotted in Fig. 5. Starting from the raw TomoSAR point clouds, the outliers are filtered out using neighborhood analysis. The façade points are then extracted using point density estimation. Subsequently, each point cloud can be split into the façade part and the non-façade part. By cross-correlating their non-façade parts, the two point clouds can be coarsely coregistered within an accuracy of several meters, which gives an initial guess of the final shifting vector. The façade parts of the two point clouds are segmented into building blocks assisted by a two-dimensional (2-D) building shape layer from online map provider such as OpenStreetMap. After obtaining the segmented point clouds, L-shape is searched for each segment, and the positions of the end points of the L-shape are estimated. Therefore, the fusion of two complete point clouds are reduced to the fusion of two much sparser point clouds comprised of the virtual façade end points. Thus, the same random sample consensus (RANSAC) point correspondence matching procedures introduced in [1] can be applied. The final shifting vectors are estimated using (3).

### A. Point Cloud Filtering

As mentioned in Section I-C, the overall geolocalization quality of a raw TomoSAR point cloud is subject to the high

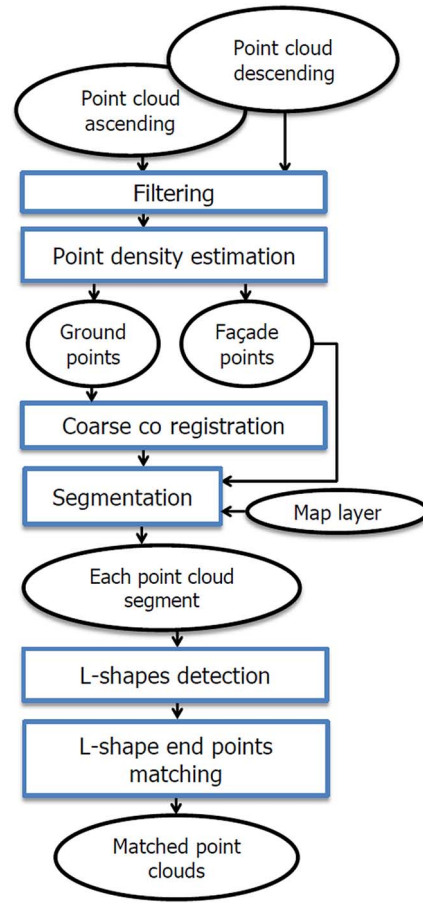


Fig. 5. Flowchart of the proposed algorithm. The black ellipses represent input/output data. The blue rectangles are procedures/processes.

dynamic range of the SAR image pixels' SNR. The criteria in model order selection for rejecting noncoherent pixels are usually set to be very soft [5], in order to preserve as much information as possible from the expensive data stacks. This leads to some amount of outliers in the resulting raw TomoSAR point clouds. Fig. 6(a) is an example of a raw TomoSAR point cloud of Berlin central station. Outliers appear as single point or small clusters elevated from the main cluster nearby. They usually have less neighboring points and experiencing larger distances to their nearest neighbor.

Therefore, the outlier filtering can be done based on the neighborhood analysis of each point in the raw point cloud. We use the mean distance to  $k$ -nearest neighbors as the single criteria, i.e., for each point, we look for  $k$  points that are closest to it, and calculate the mean distance. Points with mean distance larger than a certain threshold are discarded. For instance, in Fig. 7(a), the mean distances of the points in Fig. 6(a) to their 20 nearest neighbors are plotted as a top view. The mean distance is color coded according to the color bar. Closely clustered points have small distances showing in blue, while outliers show larger distances and are usually displayed in red. Fig. 7(b) is the histogram of the  $\log_{10}$  of this mean distance. The histogram shows two major distributions: 1) the left one refers to the main structures, and 2) the right one corresponds to points away from the main structures. The outliers can be removed

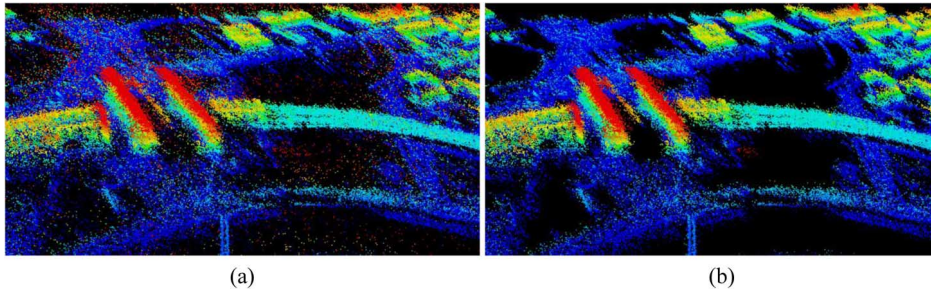


Fig. 6. (a) Raw point cloud of Berlin central station and (b) filtered point cloud. Color represents the height.

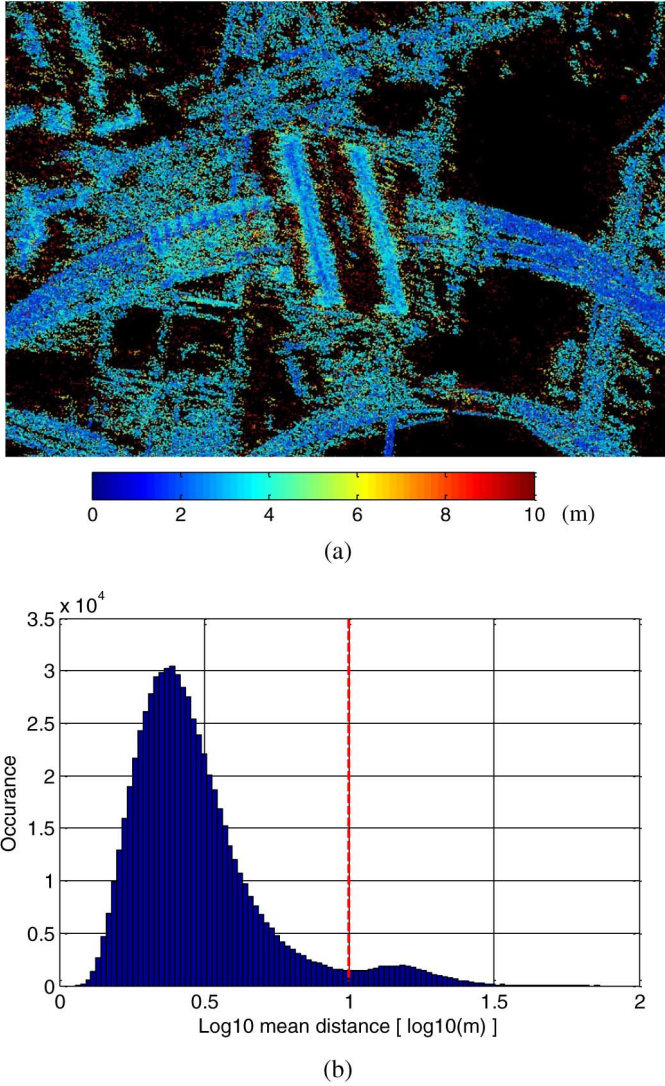


Fig. 7. (a) Mean distances of the points in Fig. 6(a) to their 20 nearest neighbors and (b) the histogram of the log 10 of this mean distance. The red dashed line indicates the cutoff distance which is 10 m.

by thresholding (indicated by the red dashed line) on the mean distance. As shown in the filtered point cloud in Fig. 6(b), most of the outliers are effectively removed. We found a threshold at 10 m and using 20 nearest neighbors are effective for TomoSAR point cloud derived from TerraSAR-X high-resolution spotlight image stacks.

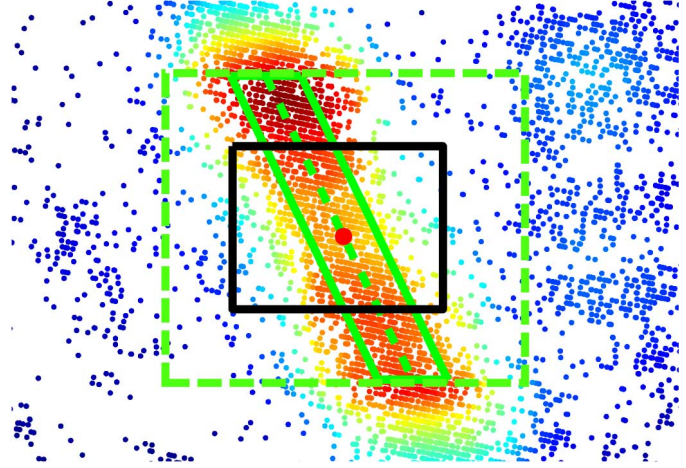


Fig. 8. Schematic drawing of point density estimation using directional window. The colored points are the projection of the 3-D point cloud in *east-north* plane, with the color indicating the point density. Blue indicates low density and red indicates high density. The color bar is not explicitly given for simplicity. The green-dashed line segment is a robust fit to the 2-D projection in a selected area. Points within the region bounded by the solid green lines are considered in the directional point density estimation, while those within the black box are considered in the box-car density estimation.

### B. Point Density Estimation Using Directional Window

The point density per square meter is employed for façade point detection. Because of the vertical accumulation of the points on the façade, their point density is usually much higher over other areas. In order to better distinguish the façade points from other, we introduce a point density estimation method using an adaptive directional window, instead a moving box.

The approach is illustrated in Fig. 8. It shows the projection of a point cloud in the horizontal plane, with the color indicating the point density, and the red dot being the target point whose point density is going to be estimated. The estimation follows three steps: 1) points within an area (the green-dashed rectangle) centered at the target point is selected; 2) a straight line segment (the green-dashed line) passing through the target point is fitted using robust algorithm, e.g., RANSAC [47], robust principle component analysis [48], etc.; and 3) only the inliers (within the solid green parallelogram) of the line segment are counted for the point density, instead of counting all the points in the black rectangle.

For comparison, Fig. 9(a) and (b) shows the point density result using directional and rectangle window, respectively, for



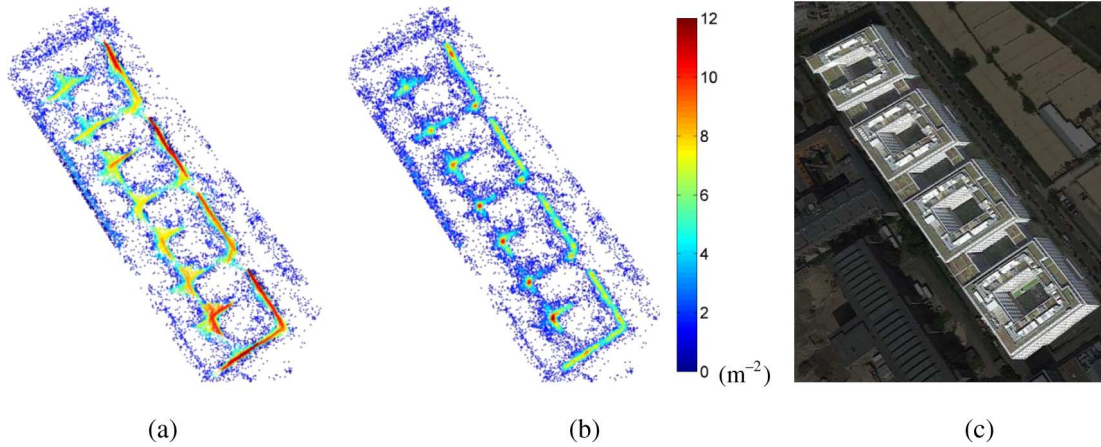


Fig. 9. Comparison of point density estimated using (a) directional window and (b) rectangle window. The color bar applies for both (a) and (b). For reference, (c) is the optical image of the same area from Google. The area of directional and rectangle windows are identical.

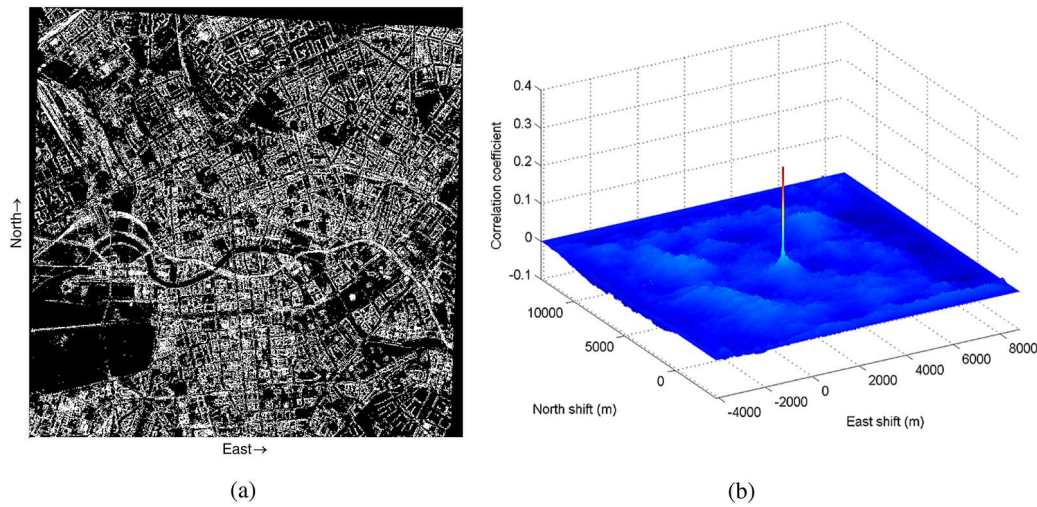


Fig. 10. (a) Binary image of the 2-D footprint of ascending point cloud of Berlin and (b) the 2-D cross correlation coefficient of the footprints of ascending and descending point clouds. The position of the peak value is the estimated coarse shift between the point cloud and the map layer.

a test area near Berlin central station. Since the same color bar is applied to the two figures, it is easy to tell that the directional window emphasizes the façade point, meanwhile keeps the point density of non-façade points unchanged compared with rectangle window. Since the urban buildings mostly consist of linear façades, our method works quite well in emphasizing façade points. The final façade points are selected by thresholding on the point density.

### C. Coarse Coregistration

The point clouds need to be coarsely coregistered to obtain an initial solution. This is a common practice in point cloud coregistration, which limits the search space in the refine coregistration step.

The coarse 3-D coregistration is done separately in horizontal and vertical directions. Horizontally, we cross-correlate the 2-D footprints of the non-façade parts of the two point clouds; and vertically, by simply aligning the mean height of the two point clouds. The 2-D footprints are rasterized binary images with a sampling spacing of a few meters. For example, Fig. 10(a) is

the binary image of the ascending point cloud of Berlin. The descending one will look similar. Their 2-D cross-correlation coefficient is in Fig. 10(b), with brighter pixel being higher correlation. The position of the peak value is the coarse shift amount.

### D. Point Cloud Segmentation

Since L-shape is a local feature of each building, the two point clouds are segmented to building blocks. Since the segmentation only needs to distinguish different building blocks, and it is also not the main focus of the algorithm, we make use of the building shape layer [49] from OpenStreetMap for assistance.

The building shape layer is a binary mask of the buildings [see Fig. 11(a)]. It is naturally not aligned with the point clouds footprints due to the shifted position of the point clouds, and sometimes the incorrect geolocation of the map itself. The alignment of the building shape layer with the point clouds 2-D footprint is done automatically using the same 2-D cross correlating technique explained in the previous section. After the

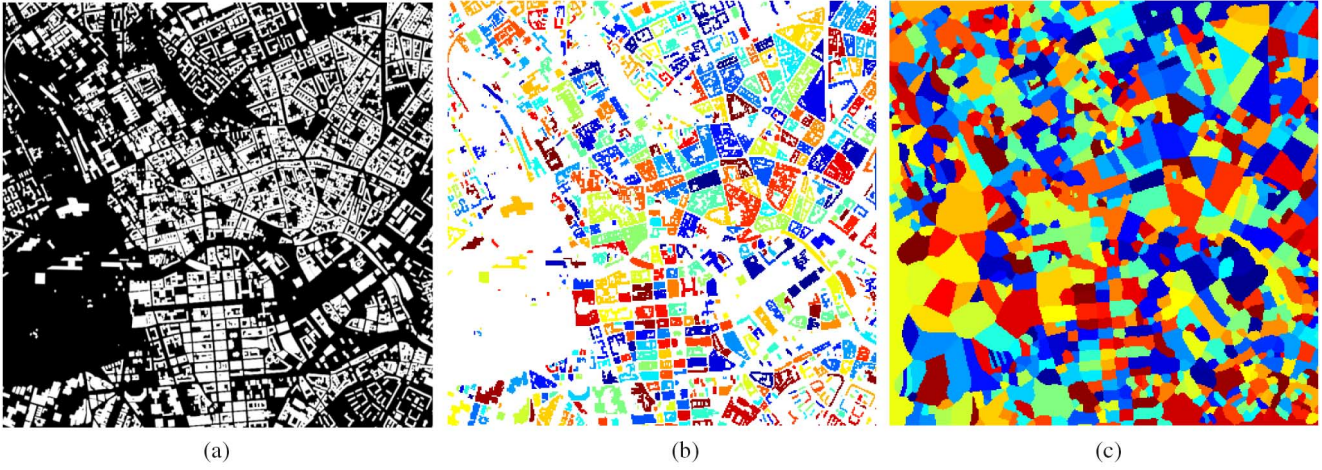


Fig. 11. (a) Building shape layer extracted from OpenStreetMap, (b) labeled individual buildings, and (c) segmentation mask derived from (b). The color assigned to each segment is random.

alignment, each building in the mask is labeled by checking their connectivity, yielding Fig. 11(b), where a random color is assigned to each building for better visualization. The unlabeled areas, i.e., the white region in Fig. 11(b), are assigned to their nearest building, giving the final segmentation mask in Fig. 11(c).

#### E. L-Shape Detection Using Weighted Hough Transform

For each point cloud segment, the façade part is selected, and one or zero L-shape is detected from its projection in the *east-north* plane. The detection is achieved by catching two interconnected line segments using weighted Hough transform. The proposed weighted Hough transform sums up the density of the points inside a Hough bin  $\Delta\theta$  and  $\Delta r$ , instead of only counting the number of points, i.e.,  $H = \sum_{i=1}^N d_i$ , with  $H$  being the Hough transform value,  $N$  the number of points in the Hough bin, and  $d_i$  the point density of each point. This concept is also illustrated in Fig. 12, where the green lines bound the Hough bin, and the size of the points inside the Hough bin denotes their point density. Both  $\Delta\theta$  and  $\Delta r$  are exaggerated for visualization purpose. Standard Hough transform gives the same value for the two bins, but the proposed weighted Hough transform favors the right bin, since its point density is generally higher.

After the weighted Hough transform, the brightest pixel in the Hough matrix is first extracted. Its corresponding line is the most prominent one in the point cloud segment. It is taken as the first line of the L-shape. The neighborhood value of this pixel is suppressed in order to keep a constraint on the minimum angle of the L-shape. Among the rest, a few pixels with the highest amplitude are selected as the candidates of the second line. The second line shall be the one that connects the first one and forms the longest continuous contour. To determine the length of the two line segments, their intersection points are calculated using their equations, and the two end points are selected by locating the last point (with threshold density) passing by the lines, respectively. Nevertheless, such end points position estimation only gives a preliminary estimation of the length of the line segments. Therefore, it needs to be refined for better

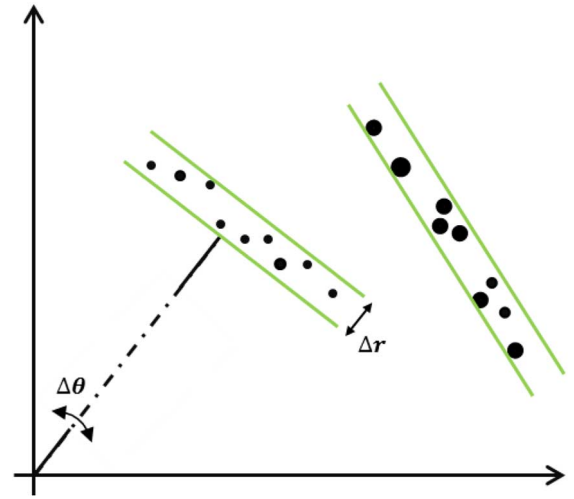


Fig. 12. Illustration of weighted Hough transform. The green arcs bounds the region of a Hough bin with spacing of  $\Delta\theta$  and  $\Delta r$ , which is exaggerated for visualization. The dots inside the Hough bins are the point cloud projection in the horizontal plane. The size of the dot represents its point density, with larger the size higher the density. Both Hough bins have nine points, which gives the same value in Hough transform. However, the right bin has higher value in weighted Hough transform.

accuracy, which is explained in the Section III. The L-shape detection is applied on each segment of the two point clouds.

Global constraints are also put on

- 1) the minimum line segment length  $l_{\min}$ , according to the knowledge of minimum façade length;
- 2) the minimum Hough value  $H_{\min}$ , which can be calculated as  $H_{\min} = \Delta r l_{\min} d_{\min}^2$ , where  $d_{\min}$  is the minimum point density of an identifiable façade in the point cloud; and
- 3) the opening direction of the L-shape depending on the orbit.

As an example, Fig. 13 shows the detected line segments and the final L-shape of the same point cloud used in Fig. 9. The detected lines are overlaid on the point density. In Fig. 13(a), the longest line segment is fixed as the first line of the L-shape, with the rest being the candidates of the second line segment.



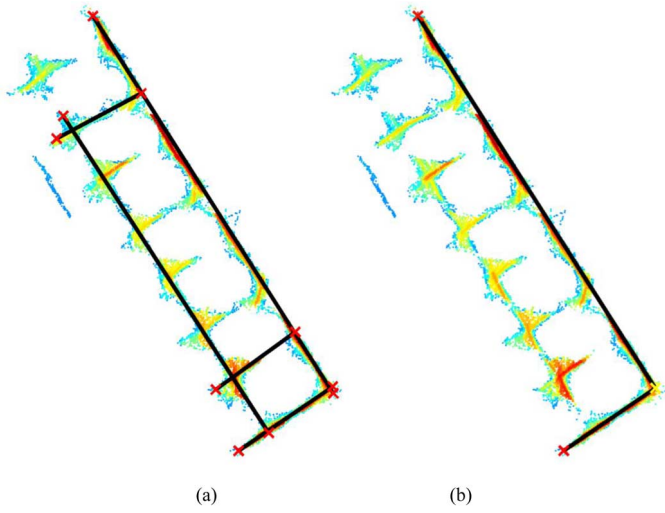


Fig. 13. Example of (a) detected line segments and (b) the final L-shape, using the same point cloud as in Fig. 9. The black lines are detected line segment. The line segments are overlaid on the point density. The roughly estimated end points are marked as red crosses. The estimation of their precise positions is explained in the next section.

Among them, the proposed algorithm automatically determines the second line by checking the connectivity with the first line and the total length of the L-shape. In the final detected L-shape, the intersection point is marked as yellow cross and the end points are marked as green.

#### F. Precise L-Shape End Points Position Estimation

The position of L-shape end point includes its horizontal 2-D coordinates and its height. The height is defined as the position where the façade intersects the ground plane, as depicted already in Fig. 4.

The horizontal coordinates are estimated using a model-based approach. Based on the detected L-shapes, the inlier points of the L-shape are projected to their corresponding line segments, i.e., façade direction. That is to say, we select only the points within a certain perpendicular distance to the L-shape and separate the two arms of the L-shape. One arm of the L-shape with inlier point should look like Fig. 14(a), where the black line is the estimated façade direction from the weighted Hough transform. For each arm of the L-shape, a projection should be obtained. The projection is basically the one-dimensional (1-D) coordinate of the façade points together with the non-façade points in the façade direction. Fig. 14(b) is the projected coordinates of the inlier points in Fig. 14(a). The middle section with much higher density belongs to the façade part. Therefore, its point density can be modeled as a rectangle function plus a constant offset for most of the urban façade.

An estimate of its point density can be obtained by convolving it with a rectangle window. For example, Fig. 14(c) shows the point density estimate of the points shown in Fig. 14(b). Its system model can be written as

$$f(x) = \int \left( A \operatorname{rect} \left( \frac{\tau - a}{2b} \right) + c \right) \operatorname{rect} \left( \frac{x - \tau}{2W} \right) d\tau \quad (4)$$

where  $f(x)$  is the estimated point density as a function of the position  $x$  in the façade direction;  $A$ ,  $a$ ,  $b$ , and  $c$  are the unknown parameter of the exact point density model, where  $A$  models the façade point density,  $a$  is the façade center position,  $b$  is the half façade length, and  $c$  is the point density of the non-façade part. Finally,  $W$  is the known half-width of the rectangle filter. Since the convolution of two rectangle functions is a trapezoid, (4) can be written as

$$f(x) = \begin{cases} 2cW & x \leq a - b - W \\ A(x - a + b + W) + 2cW & a - b - W < x \leq a - b + W \\ 2(A + c)W & a - b + W < x \leq a + b - W \\ -A(x - a - b - W) + 2cW & a + b - W < x \leq a + b + W \\ 2cW & x > a + b + W. \end{cases} \quad (5)$$

Equation (5) shows that the left and right sides of the trapezoid can fully characterize the unknown parameters  $A$ ,  $a$ , and  $b$ . Therefore, a moving fitting method is introduced to robustly fix the equation of the two sides. We use a moving window with the same width as the rectangle filter. A straight line is fitted to the points within the window. For any trapezoidal shape, high slopes should be detected at the position of the left and right sides. By detecting the position of the two high slopes, one can determine the position of the two sides. For robustness, the product of the absolute value of the slope and the number of inliers is considered in the detection of the two sides, because the maximum number of inliers should be reached when the window centered at the left or the right side. Fig. 14(d) shows the product of the absolute value of the slope and the number of inliers for the point density curve in Fig. 14(c). Two prominent peaks can be detected. The positions of the peaks are the direct estimates of the center positions of the left and right sides that are equal to  $a - b$  and  $a + b$ . Hence, they are also the estimates of the start and end position of the rectangle function that models the façade.

To ensure reliable estimation, two regularizations are performed: 1) the coarse façade length obtained from the weighted Hough transform is used as a prior to regulate the estimation; and 2) the difference in the absolute value of the two slopes is restricted, because they must be both close to  $A$  according to the model.

After obtaining the 2-D horizontal coordinates of the L-shape end point, we look for the neighboring points of the 2-D position in the ground level. The mean value of these points is an estimate of the intersection of the 2-D coordinates with the ground plane. It is taken as the height of the L-shape end point. A plane should be fitted to these points, if the local topography is not removed.

#### G. End Points Matching and Final Shift Estimation

At this step, a number of L-shape end points are found for both point clouds. Therefore, the fusion of two complete point clouds are reduced to the fusion of two much sparser 3-D point clouds comprised of only the façade end points. For robustness, RANSAC is employed to find the maximum number of matched point pairs, which is also used in [1]. More details can be found in [50]. Briefly speaking, at each iteration of



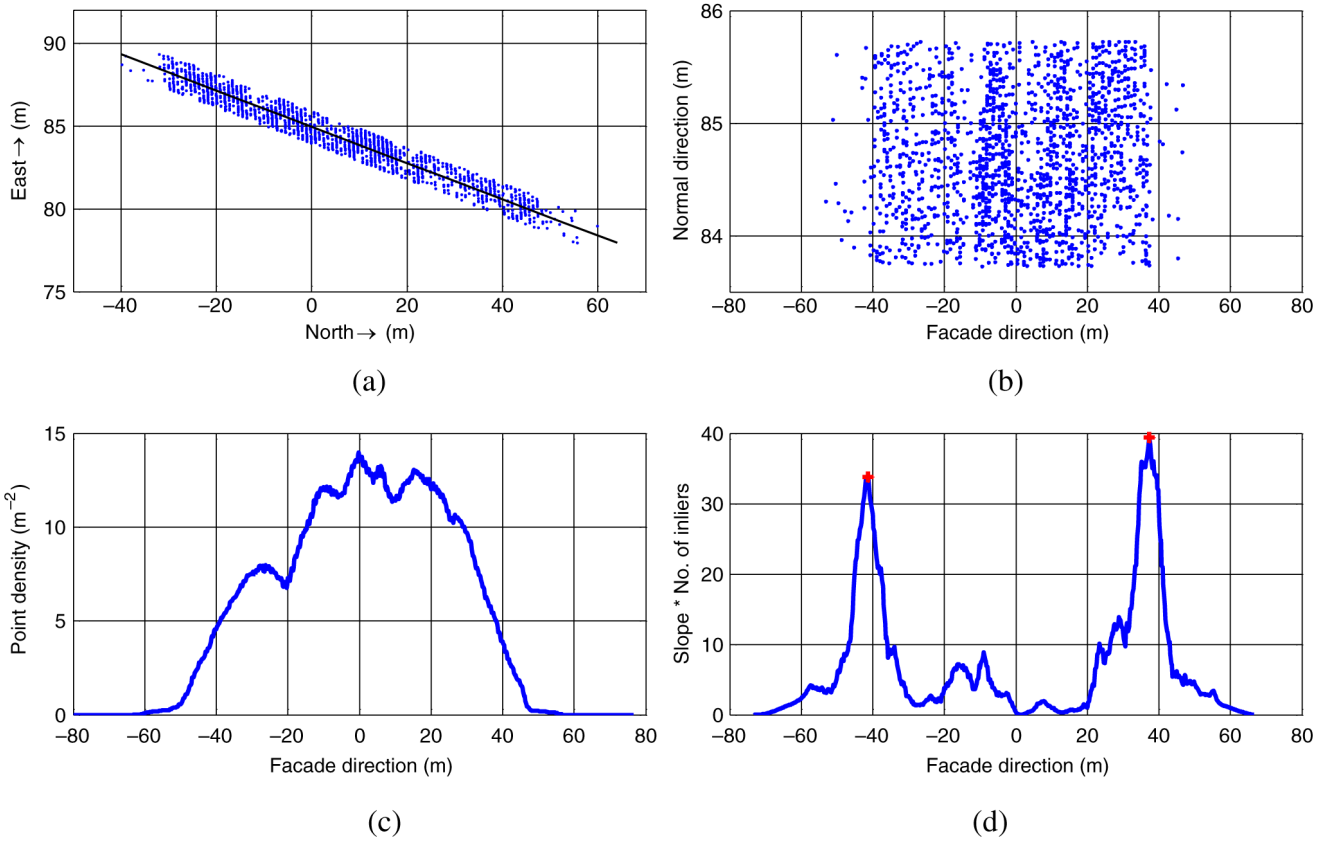


Fig. 14. (a) The inlier points of one arm of an L-shape, with the black line being the façade direction obtained from the weighted Hough transform, (b) the coordinates of the inlier points projected to the façade direction, (c) the estimate of (b)'s point density using a rectangle window, and (d) the product of the slope and the number of inliers of the fitted line using the points in the moving rectangle window along the point density estimate shown in (c). The two red crosses are the automatically detected façade start and end position at  $-41.38$  and  $37.22$  m, respectively.

RANSAC, a pair of façade end points are selected from the two sparse point clouds, least square adjustment following (3) are performed, obtaining the solution of two height offsets  $\Delta z^a$  and  $\Delta z^b$ . The two sparse point clouds are then shifted along their elevation directions. The number of matched pairs, i.e., pairs of points at close range, is then counted. The final shift is given by the one with the maximum number of matched pairs.

### III. DISCUSSION

#### A. Fusion Accuracy

The final fusion accuracy directly depends on the number of matched pairs  $N_{\text{pairs}}$  of façade end points, and the accuracy  $\sigma_{\text{point}}$  of these end point estimates, i.e.,  $\sigma_{\text{final}} = \sqrt{2}\sigma_{\text{point}} / \sqrt{N_{\text{pairs}}}$ , where  $\sqrt{2}\sigma_{\text{point}}$  is the accuracy of one pair of façade end points, according to the fusion model.

$N_{\text{pairs}}$  depends on the quality of the façade end point position estimates which are jointly affected by the accuracy of input TomoSAR point clouds, the segmentation performance, the L-shape extraction accuracy, and the number of quadrilateral buildings in the scene. According to our experience using high-resolution TerraSAR-X data, the number of quadrilateral buildings in the scene plays the most crucial role. Their typical range is a few hundreds for European cities like Berlin. We will exemplify this using real data in Section IV-B.

To analyze accuracy  $\sigma_{\text{point}}$  of the façade end points estimation algorithm explained in Section II-F, we test it using simulated data. It is generated by distributing points along the façade direction, with a constant point density on the façade part, and also a constant, but much lower point density on the non-façade part. The simulated data should look like Fig. 14(b). Important to note that the elevation accuracy  $\sigma_s$  (projected on façade direction) of the TomoSAR point cloud will affect the spatial distribution of the points. In the extreme case, when the TomoSAR point cloud is noise free, the distribution of the points should be evenly spaced in the façade and non-façade part, respectively.

We set total data length to be 40 m, façade length to be 20 m spanning from 12.00 to 32.00 m. The façade points are simulated five times, with the density being 5, 10, 15, 20, and 25/m<sup>2</sup>, respectively. The point density of non-façade area is always kept at 1/m<sup>2</sup>, and the rectangle filter size to be 5 m. Gaussian noise are then added to these point to simulate the inherent elevation accuracy  $\sigma_s$  from the TomoSAR processing, with  $\sigma_s$  being 1 and 5 m, respectively. Therefore, total 10 Monte Carlo simulations of 10 000 realizations each were performed. The simulation result is shown in Fig. 15, which shows the end point accuracy improves with respect to increasing point density. The end point location accuracy  $\sigma_{\text{point}}$  is below 30 cm in the façade direction for a typical five-story high building, i.e., with façade point density being about 15/m<sup>2</sup> in the case

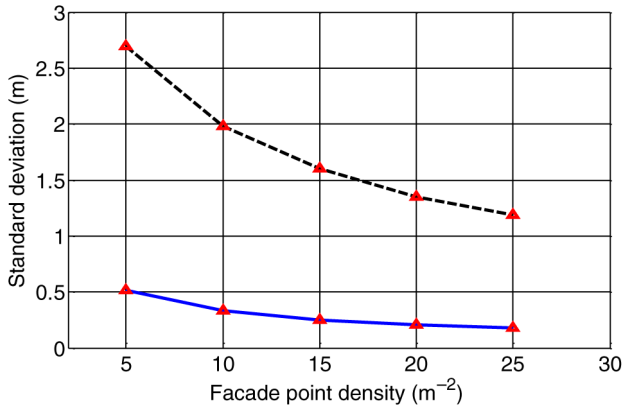


Fig. 15. Standard deviation of the end point estimates versus the façade point density. The non-façade point density is kept at  $1/\text{m}^2$ . The black-dashed line corresponds to a point cloud accuracy of 5 m in the façade direction, and the blue solid line corresponds to the accuracy of 1 m.

of meter resolution SAR data. The estimation accuracy also improves as the façade point density increases. For a typical urban area, the façade point density ranges from 5 to  $25/\text{m}^2$ , which corresponds to an accuracy of 50–20 cm. The final fusion accuracy will be typically varying from 5 to 30 cm, if we set  $N_{\text{pairs}}$  to be 100.

One should be aware of that the 3-D coordinates of a real TomoSAR point cloud are always discretized, due to the range, azimuth, and elevation sampling of 3-D SAR imaging. Therefore, the discretization in the façade direction also depends on the same facts, but in addition is subject to the heading angle of the satellite and the direction of the façade which is random. Consequently, the discretization along the façade direction is different for each façade. A simulation is performed for a discretization level of 0.5 m using the same aforementioned setting. No significant degradation of estimation accuracy is found.

### B. Parameters Settings

The proposed algorithm involves several parameters setting, including the point cloud filtering, and the weighted Hough transform for L-shape detection. Although these parameters seems to be data dependent, they are actually very much according to the inherent scale of the urban structures, e.g., the street width, number of building floors, minimum building size, etc. They can be reasonably derived with some prior knowledge of the urban structure. In the following, we give the parameter setting for TomoSAR point clouds derived from TerraSAR-X high-resolution spotlight data.

- 1) *Point cloud filtering*: we found using 20–50 nearest points, and a threshold at 10–20 m are suitable.
- 2) *2-D cross correlation*: the sampling distance of raster image should be less than half of the typical street width. We use 3 m in our algorithm.
- 3) *Point density estimation*: linear filter length is 10 m, which is a reasonable value for a shortest façade length, and width is set to be close to data resolution, i.e., 1 m.
- 4) *L-shape detection*:  $l_{\min}$  is set to 10 m, which is the shortest façade length, and  $d_{\min}$  is set to  $2/\text{m}^2$ , which



(a)



(b)

Fig. 16. (a) Google optical image of the test area and (b) the incoherent average SAR amplitude of the test area.

corresponds to the point density of a one to two-story high building in our TomoSAR point cloud. This leads to  $H_{\min} = 40$  ( $\Delta r = 1\text{m}$ ).

- 5) *Façade end points position estimation*: the rectangle filter size to be 5 m that corresponds to half of the shortest façade length.

## IV. APPLICATION ON REAL DATA

### A. Dataset

The proposed algorithm was tested on two TomoSAR point clouds generated from an ascending and a descending stack of high resolution TerraSAR-X spotlight data of Berlin. The ascending stack comprised of 79 interferograms and the descending stack has 94. The InSAR stacking and TomoSAR processing were done by the *PSI-GENESIS* [8], [51] and *Tomo-GENESIS* [18] systems, the PSI and TomoSAR processing system of Remote Sensing Technology Institute of DLR. Fig. 16 is the optical image from Google Earth, and an incoherent average of SAR amplitude of the ascending stack.

### B. Fusion Result

Each point cloud contains about 20 million points. Around 500 L-shapes are detected from each point cloud, corresponding to 1000 end points. Fig. 17(a) and (b) shows the detected L-shapes and end points of the two point clouds, overlaying on their gray-scale point density images, respectively. The matched L-shapes end points are shown in Fig. 17(c).



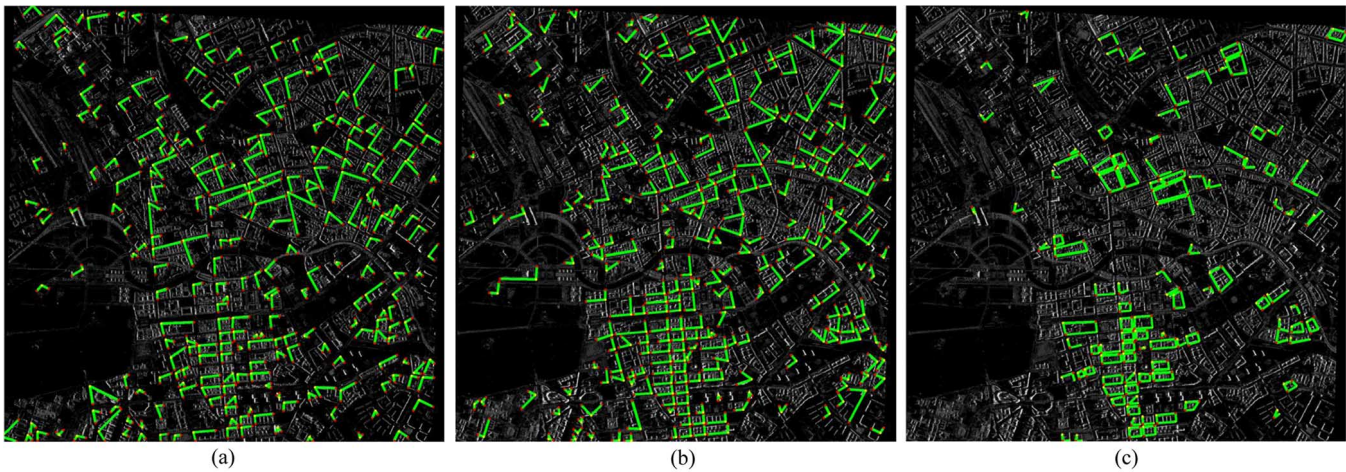


Fig. 17. (a) and (b) shows the detected L-shapes and their end points of the two input point clouds over Berlin and (c) the matched L-shapes and end points. The L-shapes are plotted in green, and the end points are marked as red. They are overlaid on the gray scale point density image.

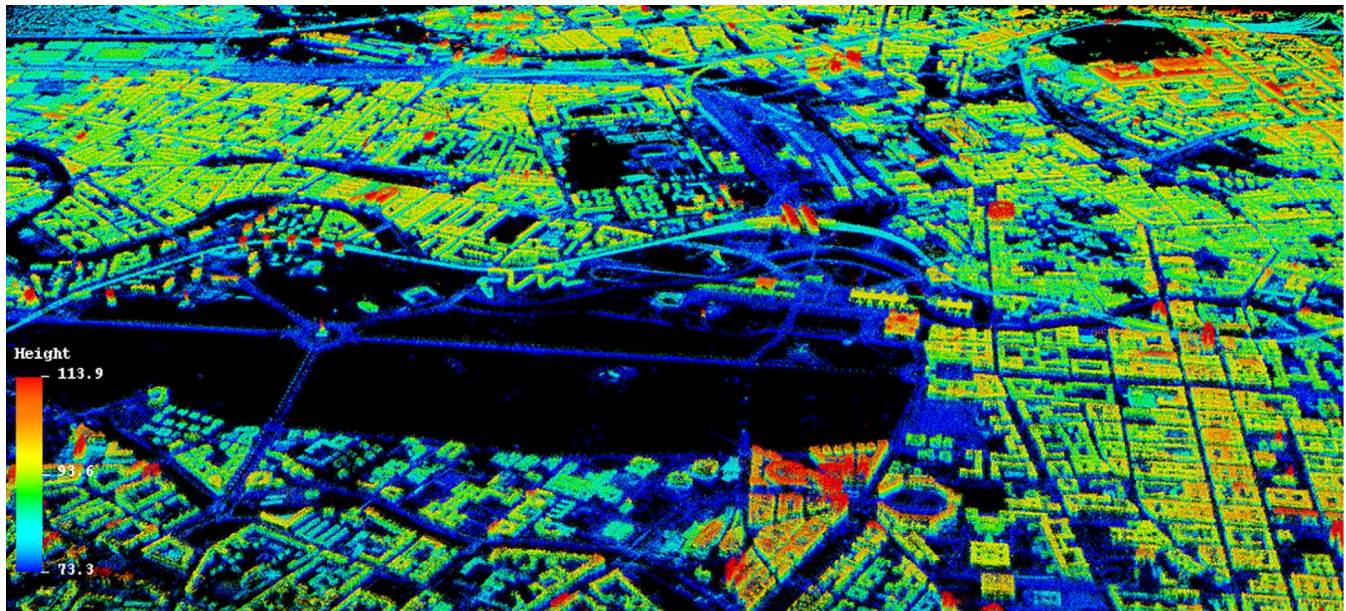


Fig. 18. Fusion result of ascending and descending point cloud of Berlin. The color indicates height on WGS84 reference surface. The unit of the color bar is meter.

Closed quadrilateral implies both pairs of end points are identified, a single L-shape implies only one pair of end points is identified. In total, 150 pairs of end points are matched, which is sufficient to robustly estimate the two unknowns  $\Delta z^a$  and  $\Delta z^b$ .

This result is compared with the result computed using the method explained in [1]. A difference of 0.51 m in *east* direction; 0.09 m in *north* direction, and almost no difference in height is found. A 3-D view of the complete fusion result is shown in Fig. 18. The height of the points is color-coded. The fused point cloud provides a Google-Street-View like visual sensation of the study area with very high level of detail. Such shadow-free TomoSAR point clouds are very useful for dynamic city model reconstruction [52] and scientific visualization [53], [54].

## V. CONCLUSION AND OUTLOOK

This paper introduced a robust method to fuse two TomoSAR point clouds derived from meter-resolution SAR data stacks of cross-heading orbits, i.e., ascending and descending orbits. The fusion is done by finding the L-shaped buildings from the two point cloud, and matching the end points of the two L-shapes belong to the same building.

The proposed algorithm was successfully tested on TomoSAR point clouds obtained from two stacks of high-resolution TerraSAR-X spotlight data. It is also compared with the fusion result using the only existing method [1] for fusing PSI point clouds derived from meter-resolution SAR data stacks of urban areas. Submeter consistency is achieved. Compared with [1], the computational speed has been greatly improved,



due to the reduced point cloud in the final RANSAC matching step. The fusion accuracy is theoretically better, since the matched pairs of points are theoretically the exact correspondence. The simulation shows the estimation accuracy of façade end point ranges from 20 cm to a few meters, depends on the façade point density (i.e., building height) and the quality of the TomoSAR point cloud itself.

The proposed method can be applied to point cloud with point density up to 10 times lower than the TerraSAR-X high resolution data, e.g., stripmap data. In order to extract sufficient number of buildings, instead of setting a threshold on point density only, a soft threshold should be set first, and then use surface normal information of the points to further classify them. This method is explained and successfully applied in [55].

The proposed algorithm requires a point cloud segmentation step, which requires either a sophisticated method or external data source, so the authors employ GIS data for assistance. The final fusion accuracy also depends on the abundance of quadrilateral buildings in the scene. Therefore, to improve the proposed method, the following work is already undergoing: 1) automatic point cloud segmentation and 2) modeling building façades using piecewise line segment instead of using L-shape only.

#### ACKNOWLEDGMENT

The authors would like to thank L. Ding from TUM-LFK for providing the source of Berlin building shape layer.

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**Yuanyuan Wang** (S'11) received his B.Eng. (Hons.) degree in electrical engineering from The Hong Kong Polytechnic University, Hung Hom, Hong Kong, in 2008, and the M.Sc. degree in Earth Oriented Space Science and Technology (ESPACE) from Technische Universität München (TUM), München, Germany, in 2010. Since January 2011, he has been pursuing the Doctoral degree with the TUM-IGSSE research team "4D City."

Currently, he is with the Helmholtz Young Investigators Group "SiPEO." In June and July of 2014, he was a Guest Scientist at the Institute of Visual Computing, ETH Zürich, Switzerland. His research interests include optimal parameters estimation in InSAR techniques, multisensor fusion algorithms of SAR/optical data, and the applications of these techniques in urban and volcanic areas.



**Xiao Xiang Zhu** (S'10–M'12–SM'14) received the Bachelor's degree in space engineering from the National University of Defense Technology (NUDT), Changsha, China, in 2006, the M.Sc., Dr.-Ing. degrees, and the "Habilitation" in the field of signal processing from Technische Universität München (TUM), Munich, Germany, in 2008, 2011, and 2013, respectively.

During October/November 2009, she was a Guest Scientist with the Italian National Research Council (CNR), Institute for Electromagnetic Sensing of the Environment (IREA), Naples, Italy. Since May 2011, she has been a Scientist with the Remote Sensing Technology Institute, German Aerospace Center (DLR), Oberpfaffenhofen, Weßling, Germany, where she is the Head of the Team Signal Analysis, and with the Chair of Remote Sensing Technology, TUM. Since September 2013, she has been leading the Helmholtz Young Investigators Group "SiPEO" and is appointed as TUM Junior Fellow. Her research interests include modern signal processing, including innovative algorithms such as compressive sensing and sparse reconstruction, with applications in the field of remote sensing such as multi/hyperspectral image analysis; advanced InSAR techniques such as high-dimensional tomographic SAR imaging and SqueeSAR; and computer vision in remote sensing including object reconstruction and multidimensional data visualization.





# Appendices

- C Wang Y, Zhu X, Zeisl B, Pollefeys M (2015b) Fusing Meter-Resolution 4-D InSAR Point Clouds and Optical Images for Semantic Urban Infrastructure Monitoring. IEEE Transactions on Geoscience and Remote Sensing. submitted.**





## Fusing Meter-Resolution 4-D InSAR Point Clouds and Optical Images for Semantic Urban Infrastructure Monitoring

Journal:	<i>Transactions on Geoscience and Remote Sensing</i>
Manuscript ID	TGRS-2015-01005
Manuscript Type:	Regular paper
Date Submitted by the Author:	28-Sep-2015
Complete List of Authors:	Wang, Yuanyuan; Technische Universität München, Lehrstuhl für Methodik der Fernerkundung Zhu, Xiaoxiang; German Aerospace Center, Remote Sensing Technology; Technical University of Munich, Signal Processing in Earth Observation Zeisl, Bernhard; ETH Zurich, Department of Computer Science Pollefreys, Marc; ETH Zurich, Department of Computer Science
Keywords:	Synthetic aperture radar, Interferometry, Image classification, Tomography, Urban areas, Image matching

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Review

# Fusing Meter-Resolution 4-D InSAR Point Clouds and Optical Images for Semantic Urban Infrastructure Monitoring

Yuanyuan Wang<sup>a</sup>, Xiao Xiang Zhu<sup>a,b,\*</sup>, Bernhard Zeisl<sup>c</sup>, Marc Pollefeys<sup>c</sup>

<sup>a</sup> Signal Processing in Earth Observation (SiPEO) Technische Universität München, Arcisstraße 21, 80333 Munich, Germany.

wang@bv.tum.de

<sup>b</sup> Remote Sensing Technology Institute (IMF), German Aerospace Center (DLR), Oberpfaffenhofen, 82234 Weßling, Germany.

xiao.zhu@dlr.de

<sup>c</sup> Institute for Visual Computing, Department of Computer Science, ETH Zurich, CH-8092 Zurich, Switzerland. zeisl@inf.ethz.ch

**KEY WORDS:** optical InSAR fusion, semantic classification, InSAR, SAR, railway monitoring, bridge monitoring

## ABSTRACT:

Synthetic aperture radar (SAR) interferometry is the only imaging-based method for assessing long-term millimetre-level deformation of individual building over a large area from space, credited to the availability of meter-resolution spaceborne SAR data, and the development in advanced InSAR techniques, such as SAR tomography. However, the inevitable SAR side-looking imaging geometry results in undesired occlusion and layover especially in urban areas, rendering the SAR images difficult to interpret.

Aiming at a semantic-level urban infrastructure monitoring, this paper proposed an algorithm to bridge the precise deformation estimates of InSAR and good visual interpretability of optical images by a strict 3-D geometric fusion of SAR and optical images, which has not been mentioned for large urban area so far. Via the precise geometrical fusion, the semantics derived from optical image can be fused to the InSAR point clouds. The proposed approach provides the first InSAR

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\* Corresponding author



point cloud of an entire urban area textured with optical attributes. Hence, the InSAR deformation analysis can be done systematically in a semantic level, instead of the current pixel-wised analysis and manual identification of the regions of interest. Examples on bridges and railway segments monitoring are demonstrated.

## 1. INTRODUCTION

### 1.1 Deformation Monitoring by SAR Interferometry

Differential synthetic aperture radar (SAR) interferometry techniques are generally recognized as the only methods so far for long-term millimetre-precision deformation monitoring over large area from space. Through modelling the interferometric phase of the scatterers, we are able to reconstruct their 3-D positions and the deformation histories. With meter-resolution SAR data, it was also demonstrated that even individual building could be monitored in very high level of detail from space.

Persistent scatterer interferometry (PSI) [1]–[4] is the workhorse among the SAR interferometry (InSAR) techniques, because of its computational efficiency. PSI only considers the phase of long-term stable scatterers –so called persistent scatterer (PS) – which is abundant in urban area, but can be sparse elsewhere. Distributed scatterer (DS)-based methods such as SqueeSAR and its alternatives [5]–[7] have emerged in the last few years to complement the drawback of PSI in nonurban area. They perform adaptive averaging of the DSs. Both of the above methods assume only single phase center in one range-azimuth resolution cell (i.e. one SAR image pixel). Therefore, in terms of urban area monitoring, *differential SAR tomography* (D-TomoSAR) [8]–[12] is the most competent method because of its capability of separating layovered scatterers in one pixel.

The output of InSAR methods are 4-D point cloud with 3-D position of the scatterers plus the deformation parameters, such as linear deformation rate. In general, millimetre-precision in the linear deformation rate can be achieved [2], [13].

## 1.2 Multi-view 3-D Reconstruction from Optical Images

Computer vision has been a topic of active research for decades. The development of cheap digital cameras, e.g. mobile phones, led to a wide-spread use and a large amount of image data available at hand. Reconstructing 3-D models from real-world objects or scenes has been a long-standing goal in the computer vision and photogrammetry community. As a result, many 3-D reconstruction algorithms were developed to exploit the present image data. At the core of all algorithms are local image features which are matched to each other among the set of captured images. These relations allow recovering both, the camera motion (camera parameters) and the scene structure (3-D geometry), which is why the problem is also termed *Structure-from-Motion* (SfM). Over the last decade tremendous progress has been made and led to impressive results, even modelling whole cities from optical images only [14]–[16].

The SfM algorithms mainly differ in how they reconstruct different camera poses together with the positions of the scene points after the pairwise image matching: in *incremental SfM* new views are added, in an iterative fashion by first estimating the absolute pose of a new image, and then updating of the structure [14]–[18]. Due to the incremental nature of the algorithm, it suggests itself for sequential image sequences or real-time applications. It is known to suffer from drift due to the accumulation of errors, and the success of the reconstruction is likely to depend on the image order. In contrast, recently global SfM methods consider the entire view graph at once [19]–[22]. Hence, they are particularly applicable to unordered image collections where no spatial or temporal order can be assumed.

If one aims to obtain a dense 3-D model, SfM is typically followed by a dense reconstructions step. They leverage the computed relative poses between cameras to estimate dense scene depth via a stereo vision algorithm such as plane sweep from two or more neighbouring views. The set of computed depth maps is then fused in a volumetric representation to obtain one consistent model [23]–[26].

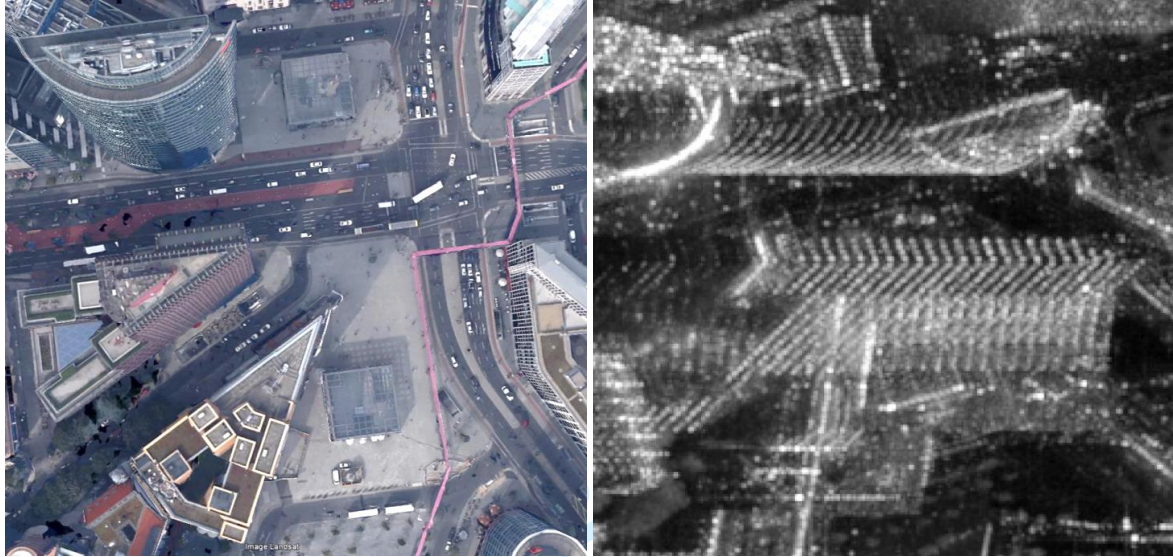
### 1.3 Motivation

The current differential InSAR (D-InSAR) methods deliver 3-D position and precise deformation estimates. However, as mentioned in [27], the inevitable SAR side-looking imaging geometry results in undesired occlusion and layover especially in urban areas, rendering the SAR images difficult to interpret. In addition, not much attention has been put on the analysis of InSAR deformation estimates after the reconstruction. The current approaches are based on manual identification of the region of interest and pixel-wised deformation analysis. It lacks a systematic way to monitor the urban area in a semantic level, e.g. systematic bridges or road network monitoring.

On the other hand, assessing long-term precise deformation using optical image SfM is not feasible at the moment. But SfM also retrieves dense 3-D position of scene points. Moreover, optical images are much easier to interpret than SAR images, e.g. visual identification of objects, perform classification, etc.

Therefore, we propose a framework for semantic urban infrastructure monitoring using a stack of SAR images and a pair of (or more) optical images. We aim to bridge the InSAR and optical field by complementing InSAR's high precision deformation estimates with the high interpretability of optical images. The semantics derived from optical images are precisely projected to the 4-D InSAR point cloud for a semantic level analysis of the deformation signal. The key lies on a strict 3-D geometric fusion of optical and SAR images. Figure 1 shows a comparison of the optical and

SAR image of a same area. The rich details in the optical image are buried underneath the strong reflection of the façade in the SAR image, causing the so-called “layover”, i.e. superposition of the reflections from multiple sources in one pixel. This calls for new fusion algorithm.



**Figure 1. Comparison of optical image (left ©Google) and SAR image (right, TerraSAR-X) of the Potsdamer platz in Berlin. The rich details in the optical image are buried underneath the strong reflection of the façade in the SAR image, causing the so-called “layover”, i.e. superposition of multiple reflection sources in one pixel.**

#### 1.4 Methodology

The general framework of the proposed approach is shown in Figure 2. The algorithm requires a stack of SAR images and a pair of (or more) optical images. The procedures are as follows.

- a. Retrieve the 3-D positions and deformation parameters of the scatterers from the SAR image stacks. Since urban area is of our main interest, D-TomoSAR is employed in order to resolve a substantial amount of layovered scatterers.
- b. Retrieve the 3-D positions of scene points from the optical images using SfM. For covering

large urban area, aerial or spaceborne images are preferred. This step also calibrates the camera parameters.

- c. Geometrically co-register the InSAR point cloud and the optical point cloud.
- d. Texture the InSAR 4-D point cloud with attributes derived from optical images, e.g. RGB colour, classification label, etc.
- e. Perform further analysis on semantic-level in the InSAR point cloud based on their semantic class.

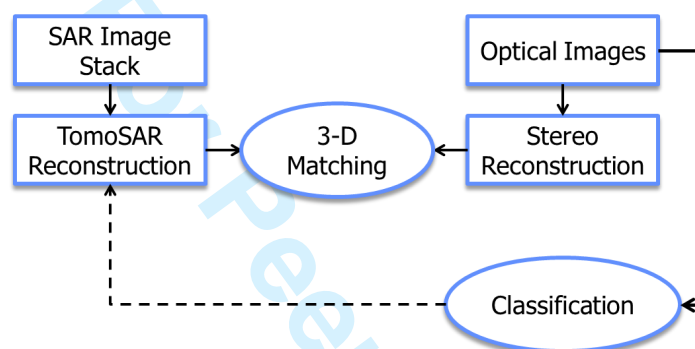


Figure 2. Flowchart of the proposed method. The semantic classification of the InSAR point cloud is achieved by co-registering the InSAR point cloud and the optical image to a reference model.

If only a single optical image is available, one should try to match higher level features, e.g. lines [28], in the optical image and TomoSAR point cloud.

### 1.5 Related work on optical and SAR image fusion

There have been many work regarding the fusion of optical and SAR images. The fusion follows basically three categories: 1) geometric fusion, 2) joint height reconstruction, and 3) joint classification.

#### Geometric co-registration



Geometric fusion is the basis of most of the higher level fusion algorithms. The current techniques are mostly based on matching of 2-D spatial features, e.g. contours, edges, templates, etc. [29]–[34]. They are good for large scale analysis, but not applicable for the analysis of individual building, especially for high resolution data.

A precise 3-D model is crucial in co-registration of SAR and optical images, for example [35] used carefully reconstructed 3-D model from photogrammetric measurements; [36] used precise airborne LiDAR point cloud; and [37] used PS point cloud retrieved from tens of high resolution TerraSAR-X spotlight images. Without a precise 3-D model, one needs to assume a strict prior [38], such as rectangle footprint etc. Considering the complex 3-D geometry of urban area, there has not been an effective algorithm of matching SAR and optical image yet.

### Joint building height reconstruction

Several literatures also explored the potential of 3-D reconstruction using a few SAR and optical images. The joint 3-D reconstruction can be said to be contextually or geometrically. Contextually, [39] used the optical image as an *a priori* term in order to regularize the height reconstructed from radargrammetry. Geometrically, [40] projected the building footprint extracted from the optical image into the SAR image, and solved building height geometrically. Alternatively, [41] retrieved the height of building from SAR and optical images independently, and fused them together using least square adjustment. Apart from these, there are also works on building reconstruction from multi-aspect SAR images [42], [43].

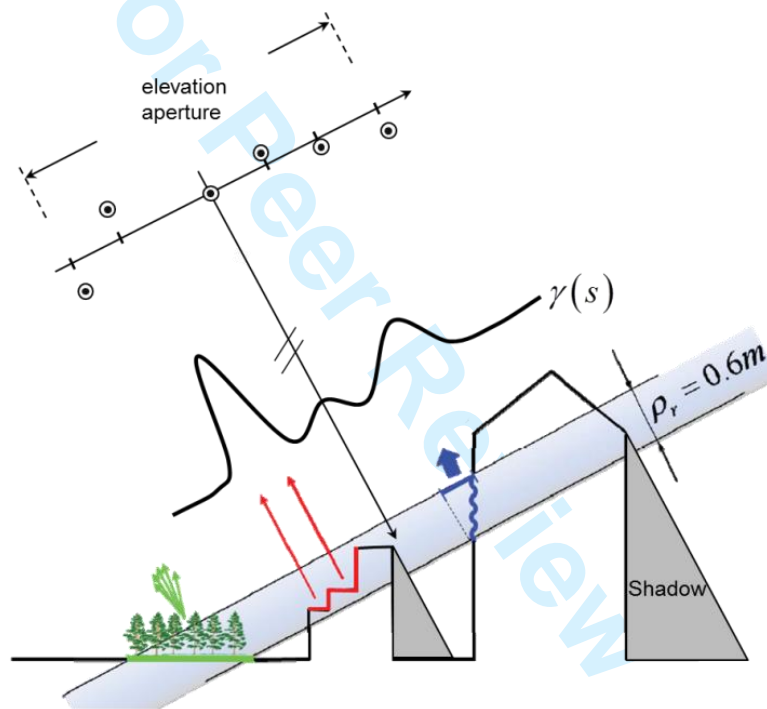
### Joint classification

Many studies have been dedicated to the joint classification of SAR and optical images. Most of the applications are on urban mapping and classification [44]–[48], since the optical and SAR image provide complimentary information in urban area. As the focus of the paper is not on the classification method, the relevant literatures are only briefly listed above for the readers' interest.

## 2. TOMOGRAPHIC SAR INVERSION

### 2.1 Principle of Tomographic SAR Inversion

The first step of the proposed algorithm is the retrieval of 4-D point clouds from a stack of multi-pass SAR images. As discussed earlier, the side-looking SAR imaging geometry leads to severe layover of scatterers and occlusion in urban area. Figure 3 shows the SAR imaging geometry in a fixed range-azimuth position, where reflections of multiple sources are layovered in one range-azimuth-elevation resolution cell.



**Figure 3. SAR imaging geometry (modified after [12]). Multiple scatterers are layovered in one resolution cell, which became one range-azimuth pixel. TomoSAR retrieves the whole reflectivity profile  $\gamma(s)$  using multiple images acquired at slightly different antenna positions.**

The reflectivity profile  $\gamma(s)$  along the third dimension *elevation*  $s$  is integrated into one range-azimuth pixel, as shown in the following equation [11], [49], [50]:

$$g_n = \int_{\Delta s} \gamma(s) \exp(-j2\pi\xi_n s) ds \quad (1)$$

where  $g_n$  is the complex SAR pixel in the  $n$ th image,  $\Delta s$  is the extent of the elevation direction, and  $\xi_n = 2b_n/(\lambda R)$  is the spatial frequency which is related to the baseline  $b_n$  of the  $n$ th image, radar carrier wavelength  $\lambda$ , and the radar range  $R$ .

TomoSAR tries to retrieve the full reflectivity profile  $\gamma(s)$  from multiple co-registered observations. Equation (1) can be discretized as

$$\mathbf{g} = \mathbf{R}\boldsymbol{\gamma} + \boldsymbol{\varepsilon} \quad (2)$$

where  $\mathbf{g} \in \mathbb{C}^N$ ,  $\mathbf{R} \in \mathbb{C}^{N \times L}$  is the so-called steering matrix containing the phase of at each discrete elevation value, and  $\boldsymbol{\varepsilon} \in \mathbb{C}^N$  is the noise assumed to be complex circular Gaussian distributed. The discretization level  $L$  in the elevation is usually much larger than the number of images  $N$ . Therefore, equation (2) is often underdetermined, and can be even ill-posed due to the irregularity and small span of the baseline. Hence, regularization is required. The Tikhonov regularization of the estimator, i.e. maximum a posteriori (MAP) estimator, is [51]

$$\hat{\boldsymbol{\gamma}}_{MAP} = \min_{\boldsymbol{\gamma}} \|\mathbf{g} - \mathbf{R}\boldsymbol{\gamma}\|_2^2 + \alpha \|\boldsymbol{\gamma}\|_2^2 \quad (3)$$

where  $\alpha$  is the regularization parameter. We realized the MAP by applying Wiener weighting on the singular values of  $\mathbf{R}$  [12]. And hence, the algorithm was named *SVD-Wiener*.

From the reconstructed reflectivity profile, multiple scatterers are separated, and hence the full 3-D (range, azimuth, and elevation) reflectivity distribution is obtained. TomoSAR is the strictest way of 3-D SAR imaging. The motion of scatterers can also be modelled and estimated by introducing new dimensions into the TomoSAR system model attributing to the motion of the scatterers [10]. For a

motion model with  $M$  parameters, the D-TomoSAR system model is a  $M+1$  dimensional discrete Fourier transform [52]:

$$g_n = \int_{\Delta p_M} \cdots \int_{\Delta p_1} \int_{\Delta s} \gamma(s) \delta(p_1 - P_1(s), \cdots, p_M - P_M(s)) \exp(-j2\pi(\xi_n s + \eta_{1,n} p_1 + \cdots + \eta_{M,n} p_M)) ds dp_1 \cdots dp_M \quad (4)$$

where  $p_1, \cdots, p_M$  and  $\eta_{1,n}, \cdots, \eta_{M,n}$  are the  $M$  motion parameters and their corresponding temporal frequencies. For example,  $\eta_n = 2t_n/\lambda$  for linear motion model and  $\eta_n = 2\sin(t_n - t_0)/\lambda$  for seasonal motion model, with  $t_n$  being the acquisition time of the  $n$ th image w.r.t. the master image.

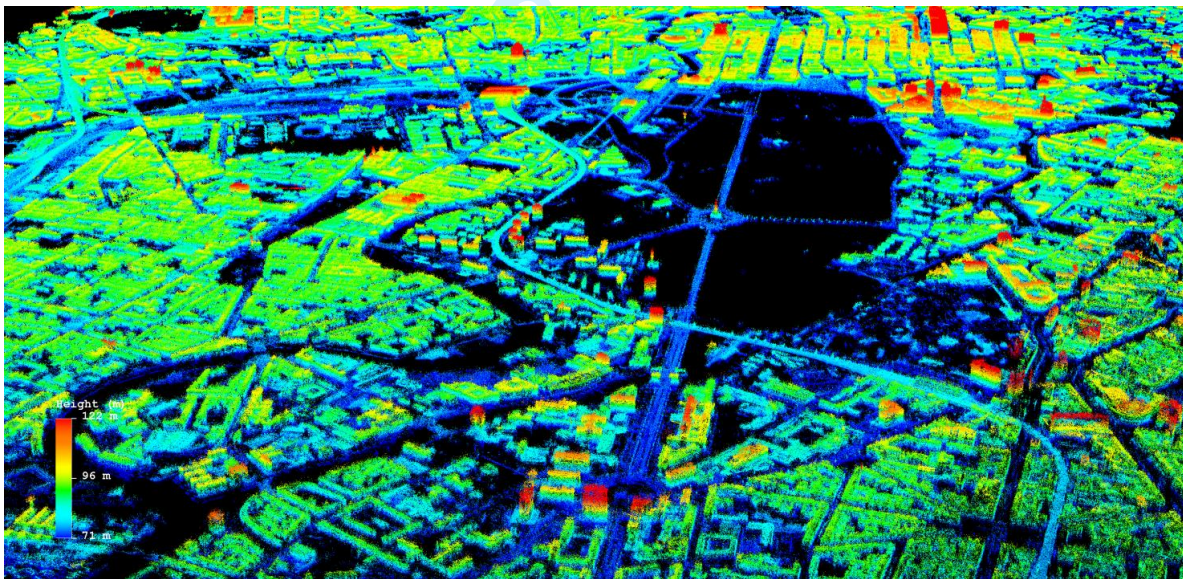
## 2.2 D-TomoSAR of Large Urban Area

The 3-D TomoSAR point clouds derived from meter-resolution SAR image stacks are the basis of the work of this paper. As D-TomoSAR is the most computational expensive multi-baseline InSAR technique, we have implemented our algorithms in an integrated system named *Tomo-GENESIS* [53], [54] – a parallel TomoSAR software of the German Aerospace Center (DLR) for large urban areas monitoring. It is developed based on the work of [11], [12], [52], [55]–[57]. It delivers, for the first time, a 5-D point cloud – 3-D position plus linear deformation rate and amplitude of seasonal motion - from a stack of SAR images. The point density is up to  $10^6/\text{km}^2$  when using TerraSAR-X high resolution spotlight data. Such density is comparable to some light detection and ranging (LiDAR) product, and enables the retrieval of the features of individual building.

We performed the D-TomoSAR reconstruction of two stacks of TerraSAR-X high resolution spotlight images acquired from 2008 to 2013 with about 90 images each. The InSAR stacking was performed by the German Aerospace Center (DLR)'s *IWAP* processor—the integrated wide area

processor [58] based on its predecessor *PSI-GENESIS* [1]. The atmospheric phase correction and the following D-TomoSAR processing was performed by Tomo-GENESIS.

The scatterers' 3-D positions were geo-coded into Universal Transverse Mercator (UTM) coordinate, so that the results from multiple data stacks with different viewing angles can be combined to fill the shadowed scene parts. For our test area Berlin, two TerraSAR-X high resolution image stacks – one ascending orbit, the other descending orbit – were processed. These two point clouds are fused to a single one, following a feature-based matching algorithm which estimates and matches common building edges in the two point clouds [59]. The combined TomoSAR point cloud has about 50 million points. Behind each of these points, the linear deformation rate and the amplitude of seasonal deformation were also estimated.



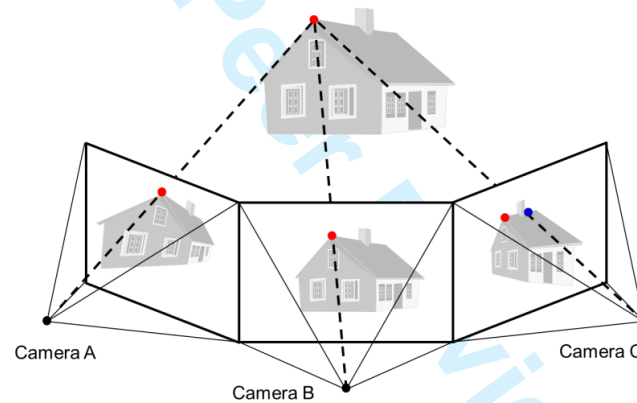
**Figure 4.** The fused TomoSAR point cloud of Berlin, which combines the result from an ascending stack and a descending stack. The height is color-coded.



### 3. OPTICAL STRUCTURE FROM MOTION

#### 3.1 Principle of Structure from Motion

The basic principle of SfM is similar to photogrammetry. It tries to reconstruct the 3-D geometry of the scene from a collection of optical images, given a set of pixel correspondence. As shown in Figure 5, if all the red dots in the image are known to correspond to the same 3-D point, one can minimize the predicted positions of the 3-D points in the images w.r.t. the observed ones, given the estimates of the 3-D points and the cameras parameters. For example, Camera C has a large reprojection error; hence, the 3-D point estimates and the camera parameters should be adjusted accordingly.



**Figure 5. 2-D projections of a 3-D house in three different camera positions and orientations.**

**The red dots are a known correspondence point between the three images. SfM estimates the unknown 3-D position of the point and the camera parameters which minimize the reprojection error (the predicted point location in image w.r.t the observed one, e.g. the blue point in Image C is the predicted location. Camera C has a large reprojection error.)**

Therefore, the SfM system model can be formulated as follows. Given a set of corresponding image points, where we use  $\tilde{\mathbf{x}}_j$  to denote the  $i$ th points in image  $j$ , the projection from their real 3-D coordinates  $\mathbf{x}_i$  is [14]

$$\tilde{\mathbf{x}}_{ij} = f_j \Pi(\mathbf{R}_j(\mathbf{x}_i - \mathbf{t}_j)) \quad (5)$$

where  $f$  is the scalar camera focal length in number of pixels,  $\mathbf{R}$ ,  $\mathbf{t}$  are the camera orientation matrix and translation vector respectively, and  $\Pi$  is the projection function:  $\Pi(x, y, z) = (x/z, y/z)$ . SfM is to estimate the unknown 3-D position, and camera parameters from the observations  $\tilde{\mathbf{x}}_{ij}$ . Assuming a Gaussian distributed reprojection noise, the maximum likelihood estimator of the parameters is [14]

$$\{\hat{\mathbf{x}}_i, \hat{\mathbf{R}}_j, \hat{\mathbf{t}}_j, \hat{f}_j\} = \arg \min_{\mathbf{x}_i, \mathbf{R}_j, \mathbf{t}_j, f_j} \sum_{i,j} \left\| \tilde{\mathbf{x}}_{ij} - f_j \Pi(\mathbf{R}_j(\mathbf{x}_i - \mathbf{t}_j)) \right\|_2^2 \quad (6)$$

Sometimes, the camera radial and tangential distortions should also be taken into account, and be estimated [17], [60].

### 3.2 Point Correspondence

Since the point correspondences are not known in prior, they must be estimated. As the estimated point correspondences are directly taken as the observation in equation (6), they become the key in the whole SfM processing.

The core of finding the correspondences is local image features [61]–[64] which are matched to each other among the set of captured images. As a consequence, the accuracy of a reconstruction depends mainly on the accuracy of local image features. One of the most successful of these features is SIFT (Scale-Invariant Feature Transformation) [63], as well as the semiglobal matching [65].

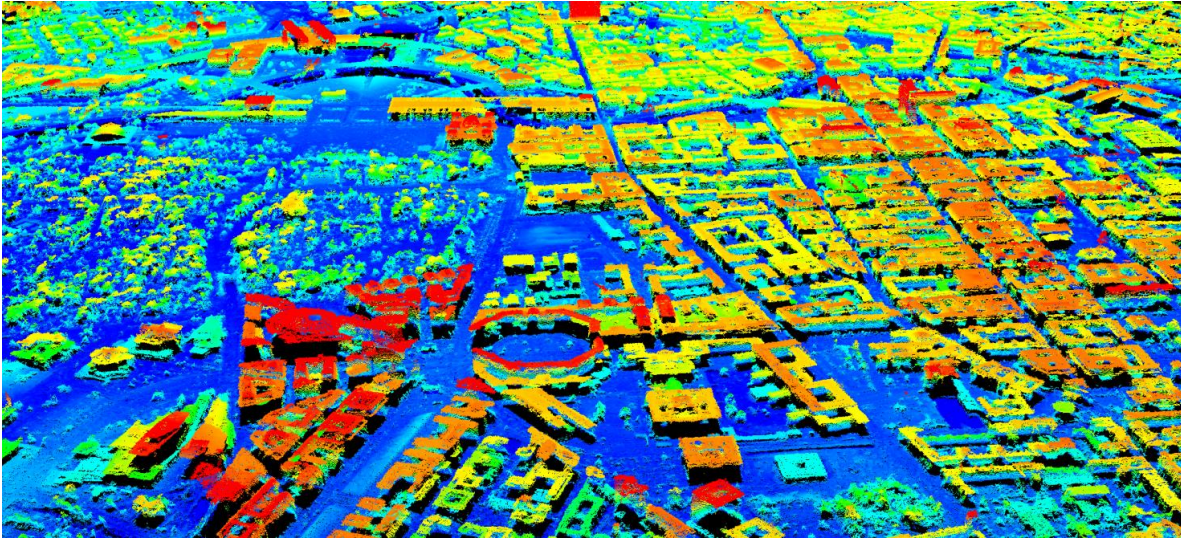
Once the features in each image are detected, they need to be matched among images. The typical approach is using the nearest neighbour search, i.e. in the feature space, finding the nearest

neighbour of a feature in one image among all the features in another image. Constrain is also put on the so-called epipolar geometry [66] to clean up the erroneous matches.

Without going too much detail, SfM can be structured into a three-step process. First, feature detection and matching are performed on a two-view basis. Second, the different camera poses together with the scene points are recovered in a global camera coordinate system. Previous calibrated two-views allow matches to be reconstructed through triangulation. Third, a global non-linear optimization, known as bundle adjustment [67], refines the estimated 3-D geometry and camera parameters.

### 3.3 Reconstruction Using ULTRACAM Data

We reconstructed the 3-D point cloud of Berlin using nine UltraCam images acquired in 2014 at an altitude of about 4000 m. The ground spacing is approximately 20cm per pixel. The initial camera positions and orientations are measured by GPS and inertial measurement unit on-board. Their standard deviation are about [35cm 35cm 20cm] for east, north, up, and  $[0.5^\circ 0.5^\circ 0.2^\circ] \cdot 10^{-3}$  for omega, phi, kappa. Therefore, a standard SfM reconstructed the whole Berlin successfully with ease. The following image is the reconstructed 3-D optical point cloud with the height being color-coded.



**Figure 6. Optical point cloud of Berlin reconstructed using structure from motion. Height is color-coded.**

#### **4. 3-D POINT CLOUDS FUSION**

Due to the relative position of the TomoSAR point cloud w.r.t. a reference point selected during the processing, it must be co-registered to the optical point cloud in order to align with the optical images.

Since both the TomoSAR and optical point clouds are geo-coded into local coordinate systems, the co-registration problem is the estimation of translation between two rigid point clouds, subject to a certain tolerance on rotation and scaling. As the optical point cloud is more accurate, it was taken as the reference.

However, a few issues should be taken into account:

- The aerial optical images we used are nadir-looking, in contrast to the side-looking geometry of SAR. In another word, façade point barely appears in optical point cloud while they are

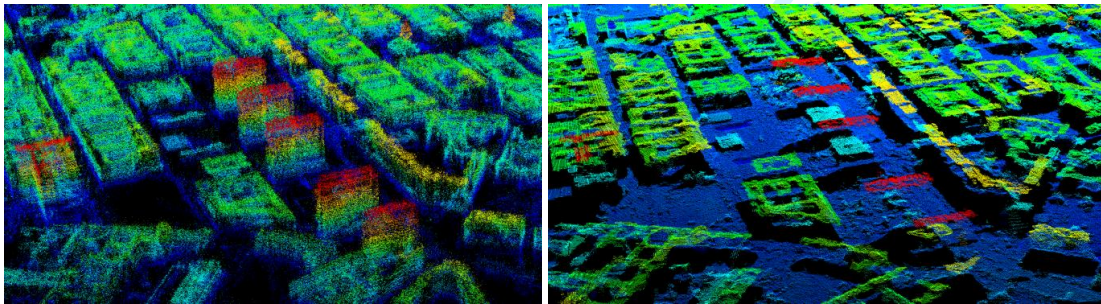
prominent in TomoSAR point cloud. This difference is exemplified in Figure 7, where the left and the right subfigures correspond to the TomoSAR and optical point clouds of the same area.

- The noise of TomoSAR point cloud is extremely anisometric. The Cram r–Rao lower bound (CRLB) of the elevation estimate for pixels of high signal-to-noise-ratio is [12], [13]

$$\sigma_s = \frac{\lambda R}{4\pi\sigma_b \sqrt{N} \sqrt{2SNR}} \quad (7)$$

where  $\sigma_b$  is the standard deviation of the baseline,  $N$  is the number of images, and  $SNR$  is the signal-to-noise-ratio in linear scale. For the SAR image stack we used, the CRLB of a 10dB pixel is about 60cm which represents almost the best situation. However, the range and azimuth are extremely precise, because of the precise orbit control of the satellite. Recent studies shows that the accuracy is up to 1cm by carefully mitigating different error sources [68]–[70]. Considering the high dynamic range of the SNR in the scene, the elevation estimates are at least one to two orders of magnitude worse than range and azimuth.

Coming to UTM coordinate, the error in elevation is mostly transferred to the east and up directions, since the satellite is roughly on a polar orbit.



(a)

(b)



Figure 7. (a) TomoSAR point cloud of high-rise buildings, and (b) the optical point cloud of the same area. Building façades are almost invisible in the optical point cloud, while it is prominent in the TomoSAR point cloud.

#### 4.1 Co-registration workflow

These unique modalities have driven our algorithm to be developed in the following way:

##### 1 Edge extraction

- a. The optical point cloud is rasterized into a 2D height image.
- b. The point density of TomoSAR point cloud is estimated on the rasterized 2D grid.
- c. The edges in the optical height image and the TomoSAR point density image are detected.

##### 2 Initial alignment

- a. Horizontally by cross-correlating the two edge images.
- b. Vertically by cross-correlating the height histogram of the two point clouds.

##### 3 Refined solution

- a. The façade points in the TomoSAR point clouds are removed, because the optical point cloud contains nearly no façade point.
- b. To refine the co-registration of the two point clouds, an *anisotropic* iterative closest point (AICP) with robustly estimated covariance matrices was introduced. The covariance is estimated using a robust M-estimator.

#### 4.2 2-D Edge Extraction

The rasterized optical height image and TomoSAR point density image are produced by computing the mean height and counting the number of points, inside each the grid cell, respectively. The edges can be extracted from these two images using an edge detector, such as Sobel filter [71]. The thresholds in the edge detector are decided adaptively, so that the numbers of edge pixels in the two

edge images are on the same order. Figure 8 is a close up view of the two edge images near downtown Berlin.

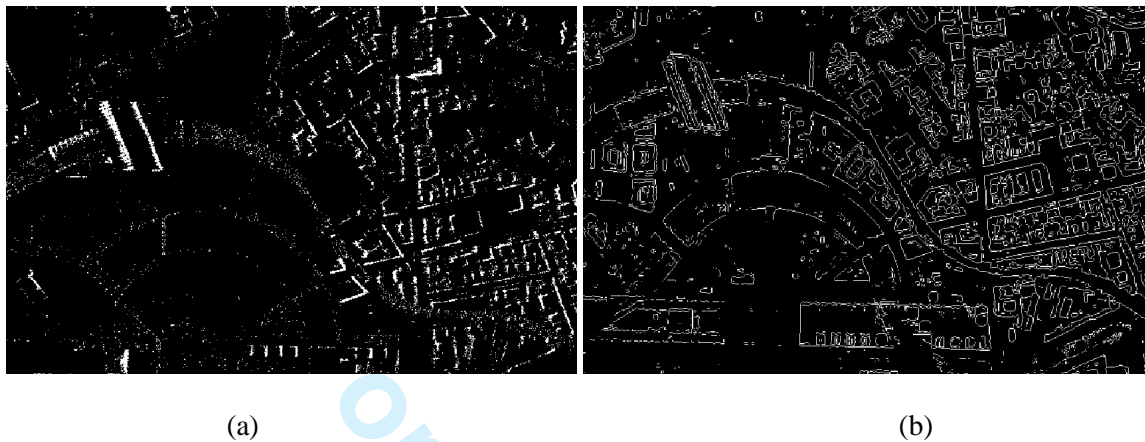


Figure 8. (a) A crop of the edge image of the TomoSAR point cloud in downtown Berlin, and (b) the edge image of the reference optical point cloud roughly at the same area.

### 4.3 Initial Alignment

The initial alignment provides an initial solution to the ICP [72] algorithm which is known to suffer from finding a local minimum.

The initial alignment in horizontal and the vertical directions are independent. The horizontal shift is found by cross-correlating the edge images of the two point clouds. A unique peak can always be found, because of the complex structures of a city. Figure 9 shows the 2-D correlation of two edge images, where a single prominent peak is found. The vertical shift is found by cross-correlating the height histogram of the two point clouds, which is shown in Figure 10.

The accuracy of the shift estimates are of course limited by the discretization in the three directions. Nevertheless, this is sufficient for our application.

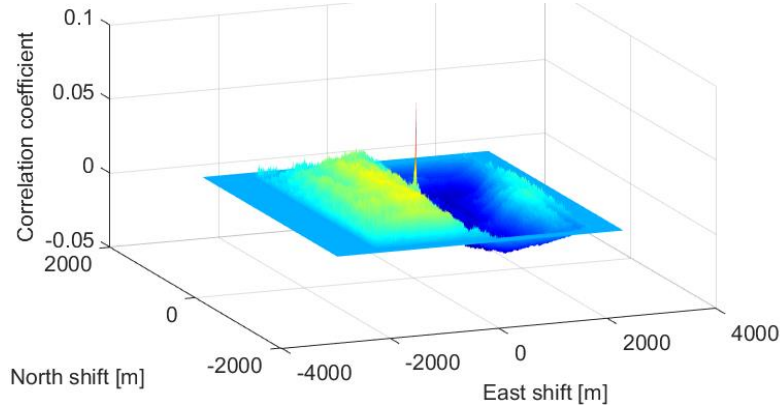


Figure 9. 2D cross-correlation of the edge images of TomoSAR and optical point clouds. Due to the pseudorandom nature of the urban infrastructure, a single prominent peak can always be found.

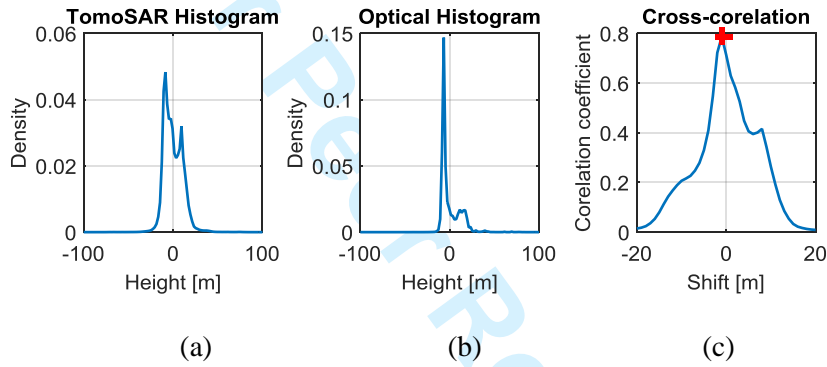


Figure 10. (a) The height histogram of TomoSAR point cloud, (b) the height histogram of optical point cloud, and (c) the correlation of (a) and (b), where the red cross marks the peak position indicating the vertical shift.

#### 4.4 Anisotropic ICP

The final solution is obtained using an anisotropic ICP algorithm based on the initial solution calculated from the previous step. Recalled the original ICP which solves the following equation

$$\{\hat{\mathbf{R}}, \hat{\mathbf{t}}\} = \arg \min_{\mathbf{R}, \mathbf{t}} \sum_i \|\mathbf{x}_i - \mathbf{R}\mathbf{p}_i - \mathbf{t}\|_2^2 \quad (8)$$

where  $\mathbf{R}$  and  $\mathbf{t}$  are the rotation matrix and the translation vector, and  $\mathbf{x}_i$  and  $\mathbf{p}_i$  are the respective corresponding points in the reference (optical) and target point clouds (TomoSAR).

Considering the anisotropic noise, [73], [74] take into account the covariance matrix  $\mathbf{C}_{\varepsilon_i \varepsilon_i}$  of the residual  $\varepsilon_i = \mathbf{x}_i - \mathbf{R}\mathbf{p}_i - \mathbf{t}$  :

$$\{\hat{\mathbf{R}}, \hat{\mathbf{t}}\} = \arg \min_{\mathbf{R}, \mathbf{t}} \sum_i (\mathbf{x}_i - \mathbf{R}\mathbf{p}_i - \mathbf{t})^T \mathbf{C}_{\varepsilon_i \varepsilon_i}^{-1} (\mathbf{x}_i - \mathbf{R}\mathbf{p}_i - \mathbf{t}) \quad (9)$$

Assuming the noise of optical point cloud is not correlated with that of the TomoSAR point cloud,  $\mathbf{C}_{\varepsilon_i \varepsilon_i}$  can be computed by

$$\mathbf{C}_{\varepsilon_i \varepsilon_i} = \mathbf{R}\mathbf{C}_{\mathbf{p}_i \mathbf{p}_i} \mathbf{R}^T + \mathbf{C}_{\mathbf{x}_i \mathbf{x}_i} \quad (10)$$

where  $\mathbf{C}_{\mathbf{p}_i \mathbf{p}_i}$  and  $\mathbf{C}_{\mathbf{x}_i \mathbf{x}_i}$  are the covariance matrix of the reference and source points, respectively.

Therefore, a correct covariance matrix estimates is the key for the anisotropic ICP algorithm. However, robustness of the covariance matrix estimation has not been addressed in any previous ICP algorithms. As mentioned earlier, the dynamic range of SAR image pixels' *SNR* are very high. Closely space pixels can have dramatic *SNR* difference, which renders their elevation estimates non-ergodic when selected for covariance estimation.

Therefore, the covariance must be robustly estimated. For this purpose, we used a robust M-estimator. Given a set of  $M$  samples  $\mathbf{X} = [\mathbf{x}_1, \mathbf{x}_2, \dots] \in \mathbb{R}$ , it is a iteratively re-weighted sample covariance matrix [75]–[77],

$$\hat{\mathbf{C}}_k = \frac{1}{M} \sum_{i=1}^M w(\varepsilon_i^T \hat{\mathbf{C}}_k^{-1} \varepsilon_i) \varepsilon_i \varepsilon_i^T \quad (11)$$

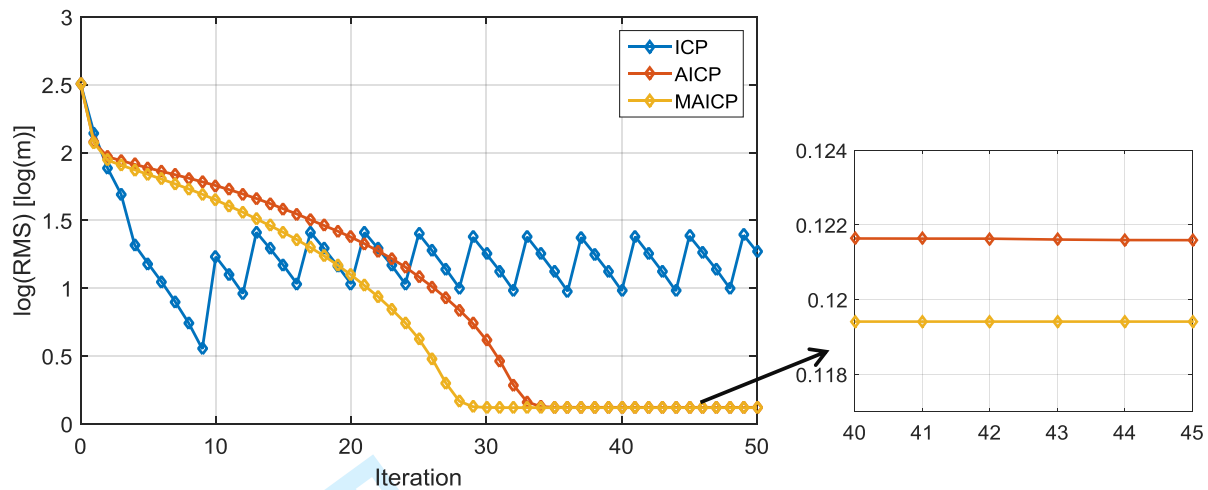
where  $k$  is the iteration index,  $\varepsilon_i = \mathbf{x}_i - \hat{\mathbf{x}}$  with  $\hat{\mathbf{x}}$  being the estimated sample mean, and  $w \in \mathbb{R}^+$  is the robust weighting function. The weighting function downweights heavily deviated samples whose whitened residual  $\varepsilon_i^T \hat{\mathbf{C}}_k^{-1} \varepsilon_i$  is large. A robust weighting function should descend to zero. We choose the t-distribution weighting function with can handle heavily tailed distributions [76]:

$$w(\varepsilon) = \frac{N + \nu}{\nu + \varepsilon} \quad (12)$$

where  $\nu \in \mathbb{R}^+$  is the degree of freedom (DoF) of the t-distribution. The lower the  $\nu$ , the more robust but less efficient (under Gaussian distributed noise) of the estimator, and *vice versa*.

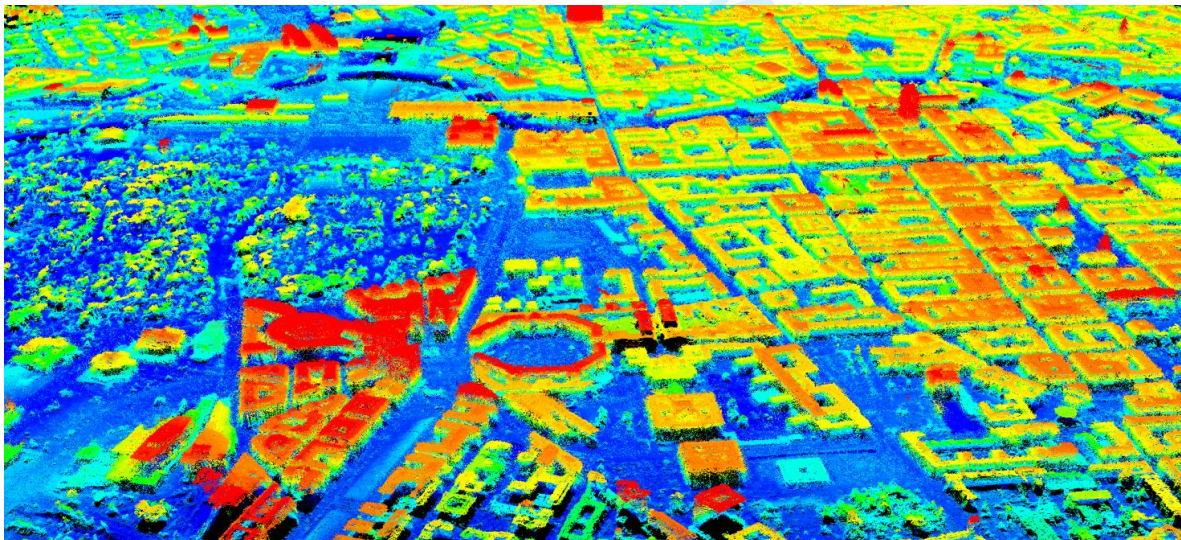
We used simulation to assess the performance of the robust AICP. A LiDAR point cloud of a building in Berlin about 10,000 points with centimetre accuracy was used as the reference point cloud. We simulated non-ergodic noise for the 10,000 points with SNR Gaussian distributed from 0 to 10dB, which well simulated the noise in our TomoSAR point cloud. The noisy LiDAR point cloud is translated to simulate the source point cloud. The following figure compares the performance of the original ICP, the AICP, and AICP with M-estimate of covariance matrix (MAICP). The figure shows the root mean square (RMS) error of the estimated at each iteration of the ICP algorithms. The original ICP suffers from finding a local minimum, and oscillates between two states. The AICP and robust AICP both converged. The robust AICP improved the convergence rate as well as the co-registration accuracy.



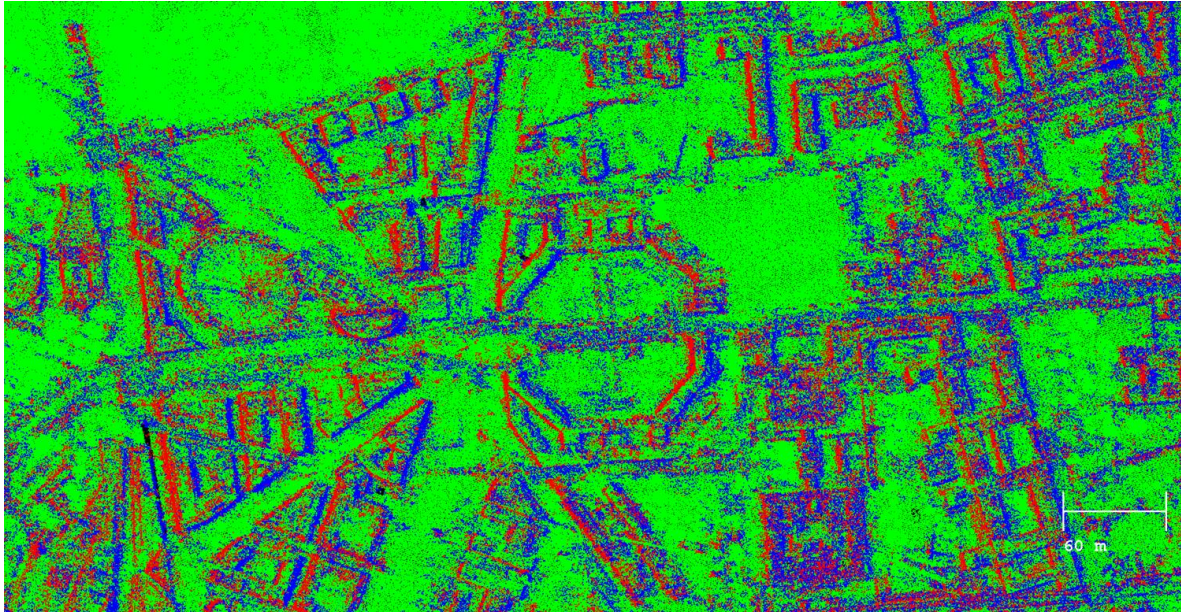


**Figure 11. Comparison of the original ICP, anisotropic ICP, and anisotropic ICP with robust covariance matrix estimate (M-estimate). The original ICP may suffer from finding a local minimum. Using a robust covariance matrix not only improves the convergence rate, but also co-registration accuracy.**

Figure 12(a) demonstrates the co-registered point cloud combining the optical images-derived one (also shown in Figure 6) and two TomoSAR point clouds (also shown in Figure 4) from ascending and descending viewing angles with color representing the height. Successful co-registration can be confirmed by seeing the correct location of the façade points in Figure 12(b) which shows the top view of fused point cloud with different colors representing different point clouds.



(a)



(b)

Figure 12. (a) The fused point cloud combining the optical point cloud (Figure 6) and two TomoSAR point clouds (Figure 4) from ascending and descending viewing angle. The height is color-coded. And (b) the top view of co-registered point cloud where green, red, blue represent the points from optical, ascending and descending TomoSAR point cloud, respectively.

## 5. TOMOSAR POINT CLOUD TEXTURING

Once the TomoSAR and optical point clouds are co-registered, the TomoSAR point cloud can be textured with attributes from the optical images, e.g. RGB color. This is also known as rendering: given a viewpoint, solve the occlusion of objects. There are two points should be taken into consideration.

- Firstly, points do not occlude each other except degenerated points, because they are dimensionless. Typically, the rendering of a point cloud requires reconstructing the 3-D surface and testing the depth of each point. This is known as the z-buffer algorithm. But reconstructing reliable 3-D surface from TomoSAR point cloud is still a challenge task, although there are



some pioneer works on reconstructing façades and roofs from spaceborne TomoSAR points [78], [79]. Therefore, the rendering technique should avoid the surface reconstruction.

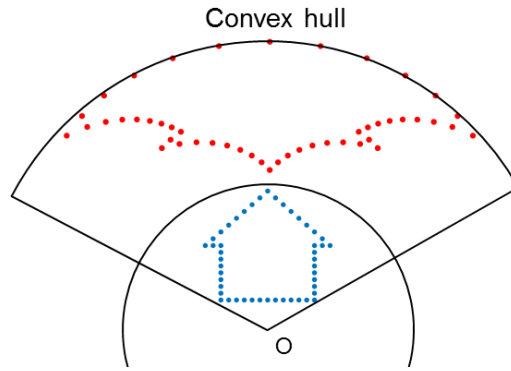
- Secondly, one point can be visible from multiple images. Strategy should be developed to choose the appropriate textures from the optical images. The textured point cloud should be visually consistent.

### 5.1 Point-based Rendering

[80] introduced a hidden point removal (HPR) operator which is an elegant solution to avoid reconstructing an underlying surface or even normals. HPR inverts the point cloud using certain function, so that all the visible points lie on the convex hull of the inverted point cloud. The most common inverting function is the *spherical flipping*. Given a point  $\mathbf{x}$  of any dimension, the spherical flipping function is defined as follows:

$$\hat{\mathbf{x}} = f(\mathbf{x}) = \mathbf{x} + 2\left(r - \|\mathbf{x}\|\right) \frac{\mathbf{x}}{\|\mathbf{x}\|} \quad (13)$$

where  $r$  is radius of the multi-dimensional sphere. The following figure illustrates the transformation function and HPR in 2-D. The blue dots are the original point cloud. And the red dots are the inverted one given a viewpoint  $O$ . The convex hull of the inverted point cloud is computed and the points lying on the convex hull are marked as visible.



**Figure 13. Hidden point removal operator. The blue dots are the original point cloud. And the red dots are the inverted one given a viewpoint  $O$ .**

However, as mentioned in [81], the original HPR performs poorly in presence of noise. Considering the much large noise in spaceborne TomoSAR point cloud than typical computer vision problem, the HPR is not directly applicable.

[81] suggested to include a buffer zone around the convex hull, so that all the point in the buffer zone can be marked as visible. Inspired by this, we introduced a *visibility probability* which is a Gaussian function of the point-to-convex hull distance  $d$ :

$$p(\hat{\mathbf{x}}) = \exp\left(-\frac{d^2}{2\sigma_{\hat{\mathbf{x}}}^2}\right) \quad (14)$$

where  $\sigma_{\hat{\mathbf{x}}}$  is the norm of the total noise standard deviations (all three directions) in the inverted point cloud.  $\sigma_{\hat{\mathbf{x}}}$  is related to original point cloud. Consider the very anisotropic noise of TomoSAR point cloud, we define the standard deviation in the native SAR coordinate range, azimuth, and elevation being  $[\sigma \quad \sigma \quad k\sigma]$ , where  $k > 1$  is a positive factor. As mentioned earlier, this factor is about 10 to 100. Hence, the noise in range and azimuth can be neglected. Transforming to UTM coordinate, the standard deviation for east (x), north (y), and up (z) is approximately

$\left[ k\sigma/\sqrt{2} \quad \sigma \quad k\sigma/\sqrt{2} \right]$  (assumed a polar orbit, and  $45^\circ$  of incidence angle). The  $\sigma_{\hat{x}}$  is derived to be:

$$\sigma_{\hat{x}} = \frac{4rk\sigma}{\|\mathbf{x}\|_{\min} - k\sigma} - k\sigma \quad (15)$$

where  $\|\mathbf{x}\|_{\min}$  is the minimum distance of the original point cloud to the viewpoint.

## 5.2 Consistency Check

The noise of TomoSAR point cloud inevitably caused some erroneous visibility estimates. In addition, many points are visible to multiple images. Points located far away from the center of the images often have oblique angle with poor resolution and optical quality. We use the information of the neighbouring points to check the global consistency of the whole rendering.

Firstly, for each point, a score similar to [28] is obtained w.r.t. each image as follows:

$$c_j = p_j d_j \quad (16)$$

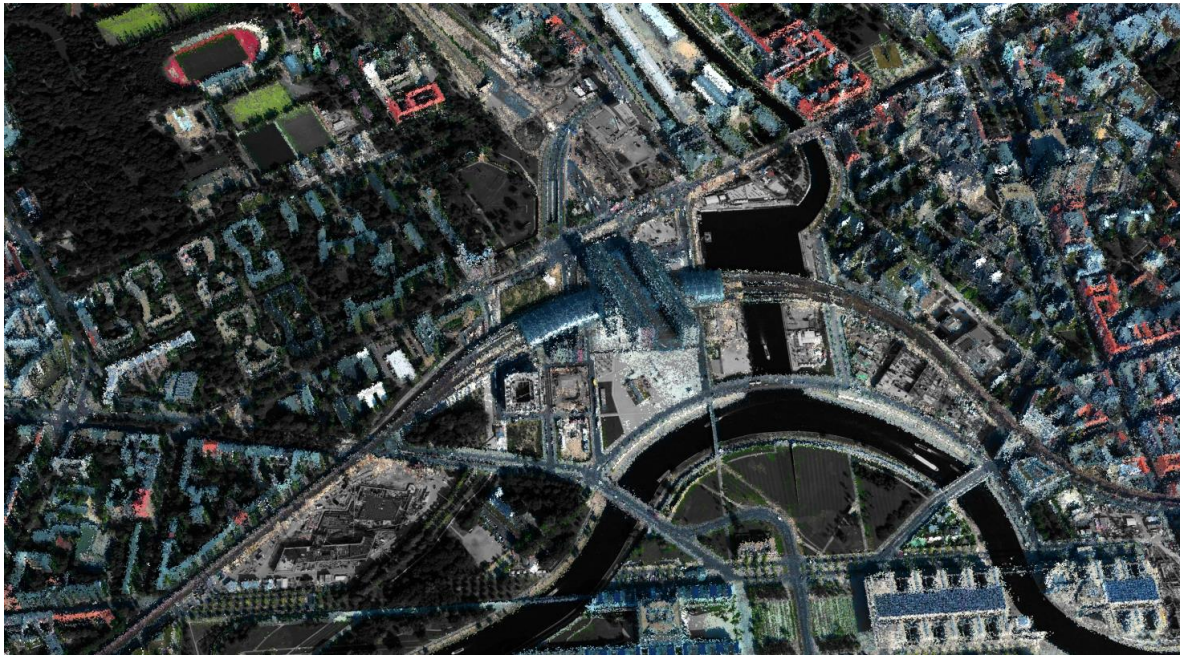
where  $j$  is the image index,  $p_j$  is the visibility probability, and  $d_j$  is the inverse of the distance to the viewpoint.  $d_j$  is regarded as the resolution of the image at the point. Equation (16) basically says image with larger visibility and shorter distance is favoured. The image with the largest score is selected as the initial texture source image.

Secondly, to decide the final texture source image, we search the  $k$  nearest neighbours of each point, and perform a weighted multi-class majority voting. The weight is a Gaussian kernel as a function of the distance. Largest weight is assigned to the point itself, and lower weights are assigned to the points further away. The image with the highest votes is selected as the final texture source.

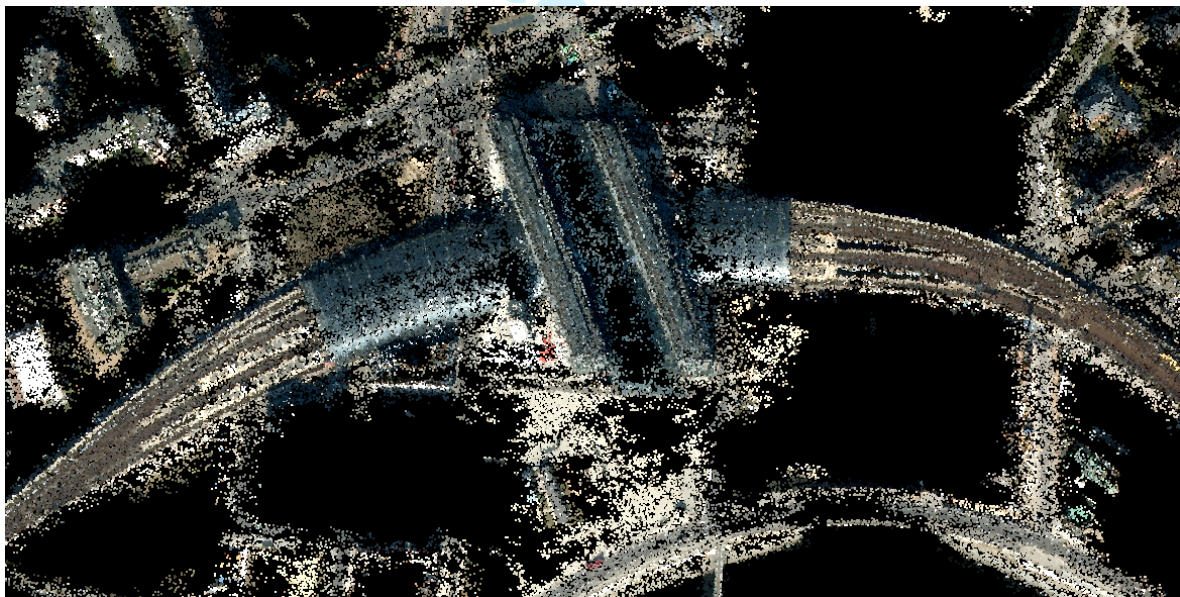


The following figure shows the first spaceborne TomoSAR point cloud textured with optical color. Where no TomoSAR point is available, the grayscale image is filled. The lower figure is a zoom in of the Berlin central station. The boundary of the railway and the building edges are precisely projected.

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(a)



(b)

Figure 14. (a) First TomoSAR point cloud textured with optical image. Color is enhanced for better visualization, and (b) zoom in to the Berlin central station. The boundary the building edge, and the railway are clearly textured.

## 6. PRACTICAL DEMONSTRATION

This section demonstrates the *semantic level urban infrastructure* by transferring the semantic classification label from the optical image to the TomoSAR point cloud. Example is shown with railway and bridge monitoring, as the subtle movement of railways and bridges can be well captured by TomoSAR, while the whole railway and bridge network can be well labelled using optical images.

### 6.1 Image Classification

The procedure of our image classification is briefly introduced, as this is not the focus of this article. Our image is classified patch-wisely using the Bag of Words (BoW) [82] method. It originates from text classification, where a text is modelled as the occurrence of the words in a dictionary, disregarding the grammar as well as the order. Analogous to text, the BoW descriptor  $\mathbf{w}$  of an image (in our case an image patch)  $\mathbf{Y}$  is the occurrence of the “visual” words in a predefined dictionary  $\mathbf{D}$ .

Define the dictionary matrix as  $\mathbf{D} \in \mathbb{R}^{N \times k}$ , where  $N$  is the dimension of the word, i.e. feature/atom, and  $k$  is the number of words. The  $k$  feature vector should include representative features appearing in the whole image, so that each patch can be well described. Therefore, the dictionary is usually overcomplete. The dictionary is usually learned from the image, by collecting representative features and then discretize into  $k$  levels.

The classification is done using a linear SVM [83] implemented in an open source library VLFeat [84].



## 6.2 Railway Monitoring

We applied the semantic classification scheme on the orthorectified image centered at the Berlin central station. It was generated from UltraCam-D images using SfM with semiglobal matching [65]. For this analysis, we particularly classified the railway and river class.

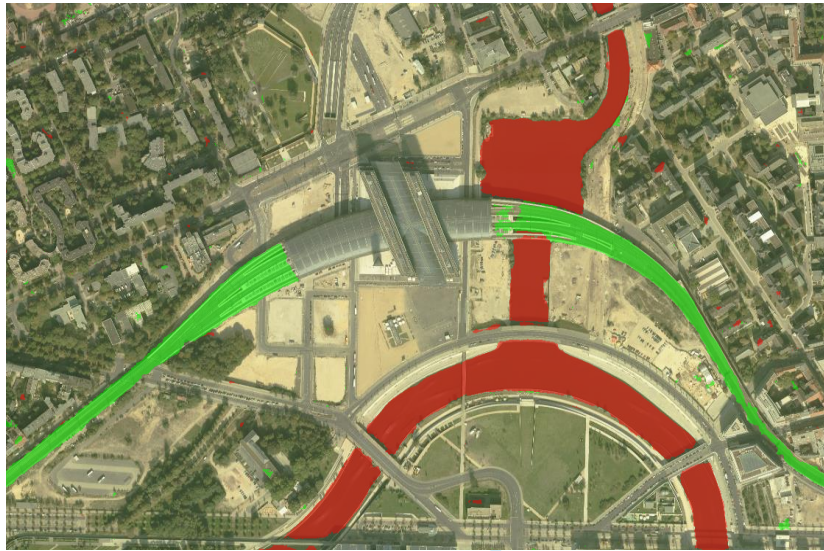


Figure 15. River (red) and railway (green) classified using the BoW method. BoW performed well. Some false alarm appeared as small clusters. They can be cleaned up using size analysis.

Based on the classification, the corresponding points in the TomoSAR point cloud can be extracted by projecting the classification label to the point cloud. A smooth spline function was fitted to the horizontal coordinates of the railway points to connect separated segments. For example, the railway label is disconnected due to the presence of the central station. Figure 16(a) shows the extracted continuous railway points overlaid on the optical image. The color shows the amplitude of seasonal motion.

This motion is most probably caused by the thermal dilation of the steel railway because of the seasonal temperature change. It can also be easily observed that the railway deformation

experiences certain discontinuity. It is suspected to be caused by the mechanical clearance between different railway segments. Therefore, it is also possible to monitor each railway segment.

However, the raw TomoSAR estimates are relatively noisy due to the relative low SNR of some scatterers. To filter out the noise, a special prior was employed, i.e. the thermal dilation of a steel beam is linearly proportional to its length at its first approximation [85]. That is to say, the railway deformation parameters is piecewise linear, and the scatterers on the same cross-section of the railway undergo similar deformation. Therefore, the deformation estimates can be filtered by minimizing its total generalized variation along the railway direction (with second order derivative which favours piecewise linear):

$$\hat{\mathbf{f}} = \arg \min_{\mathbf{f}} \left\{ \frac{1}{2} \|\mathbf{f} - \mathbf{a}\|_2^2 + \lambda \|\Delta \mathbf{f}\|_1 \right\} \quad (17)$$

where  $\mathbf{a}$  is the deformation estimates along the 1-D railway direction (i.e. stretch the railway to a straight line),  $\mathbf{f}$  is the noise-free version of  $\mathbf{a}$ , and  $\Delta \mathbf{f}$  is the second order derivative of  $\mathbf{f}$ . As shown in literatures of total generalized variation [86], [87], the  $L_1$  norm of second order derivative is convex and lower semi-continuous, one can solve it using convex optimization solvers.

The filtered deformation estimates are shown in Figure 16 (b), in parallel with the unfiltered estimates. The noise has been greatly reduced while the discontinuity is well preserved. To have a quantitative comparison, Figure 17 (a) plots the original (in blue) and the denoised (in red) deformation estimates as a function of the 1-D railway direction. The original estimates have a standard deviation about 1~2 mm which concealed some edges where the magnitude of deformation is low. The filtering successfully reconstructs the edges.



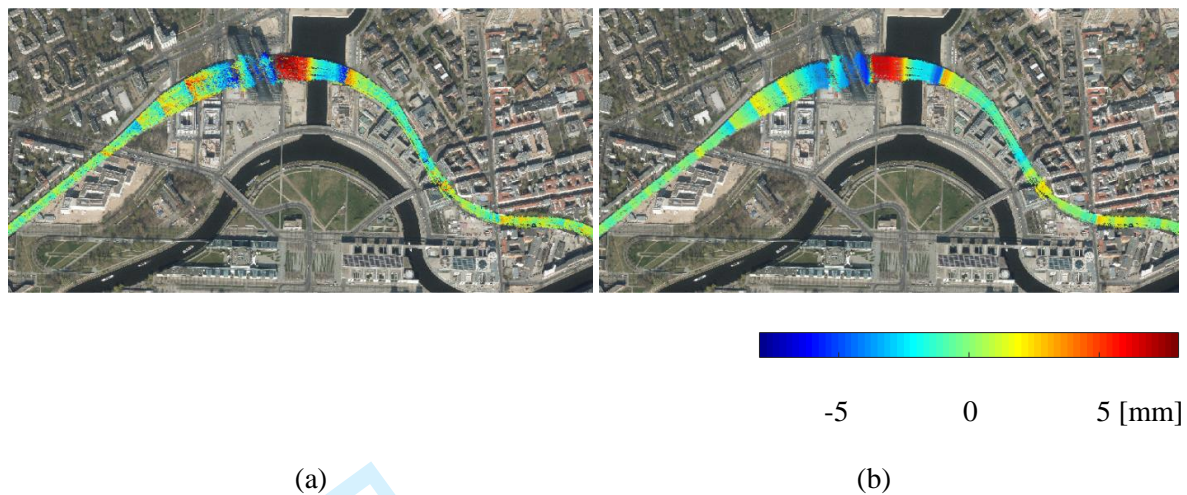


Figure 16. (a) Railway points extracted from the original TomoSAR point cloud, and (b) the railway points filtered using total generalized variation. The colour shows the amplitude of seasonal motion due to the thermal expansion of the railway.

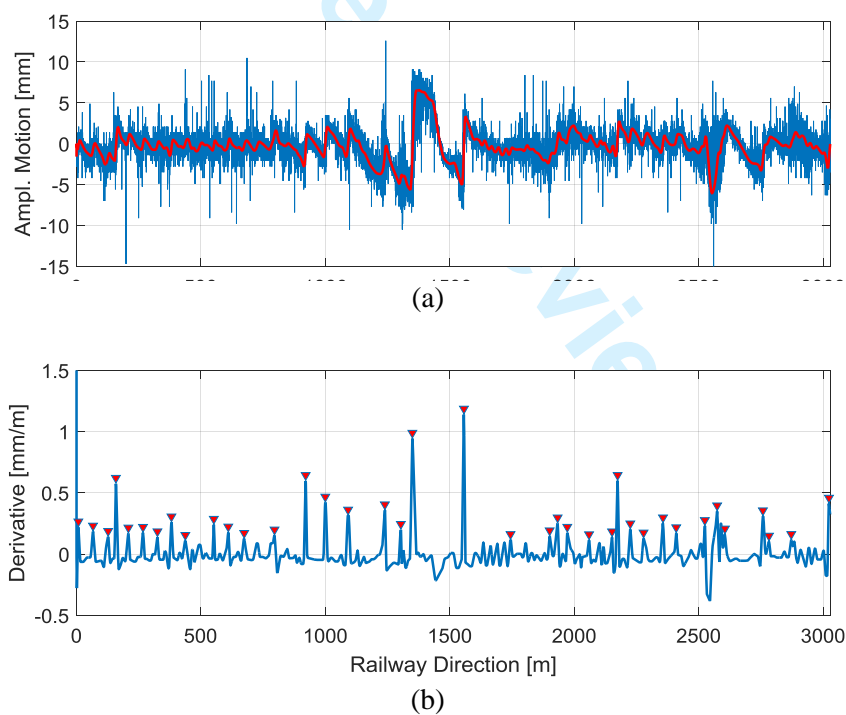


Figure 17. (a) the original and the filtered estimates of amplitude of seasonal deformation in blue and red, respectively, and (b) peaks detected in the derivative of the denoised deformation function along the railway direction. They are the locations of the railway segment joints.

By detecting the peaks in the derivative of the deformation function, the joints of railway segments can be detected. Constraint was put on the minimum distance between two peaks representing the minimum length of a railway segment. The detected peaks in the deformation's derivative can be seen in Figure 17(b). The positions of the peaks on the railway are shown as the green dots in Figure 18 upper. Each green dot represents the midpoint of the cross-section. In the lower subfigure of Figure 18, the close up view of the two joints in the optical image is provided. It can be clearly observed that the railway joint shown up as dark lines in the optical image.





Figure 18. Upper: the midpoint of the detected railway joint cross-section marked in green, and lower close up view of the railway joint. The background optical image has a ground spacing of 20cm.

### 6.3 Bridges Monitoring

By analysing the discontinuity of the river segmentation and assuming the discontinuities are caused by bridges, the bridges' positions can be detected automatically. Without going into too much detail, the extracted bridges points overlaid on the optical image are shown in Figure 19 where the color represents the amplitude of seasonal deformation. The upper most bridge belongs to a segment of the railway which is known to have thermal expansion. The middle bridge undergoes a 5mm seasonal motion on its west end and 2mm at the east end. This suggests a more rigid connection of the bridge with the foundation at its east end. The two lower bridges are stable according to the motion estimates.

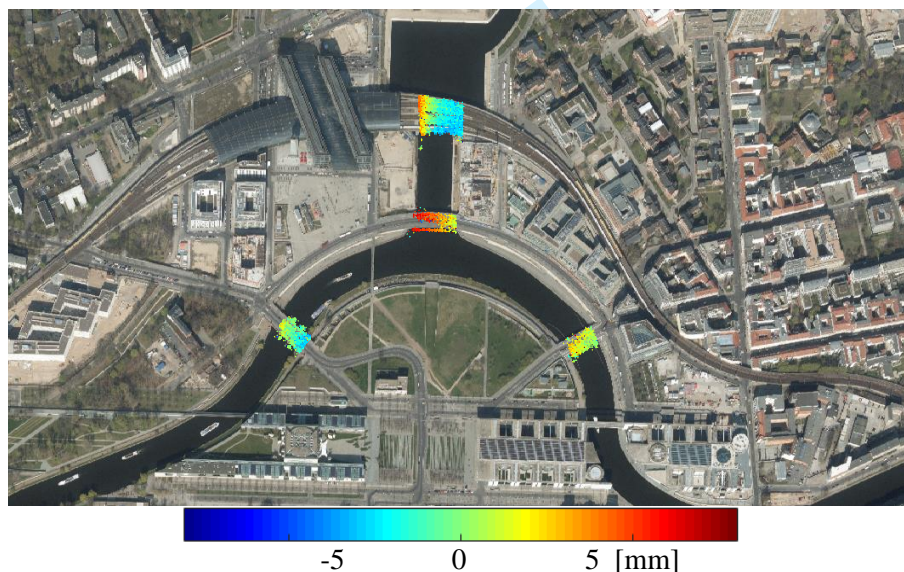


Figure 19. Overlay of the amplitude of seasonal motion of bridges extracted from the TomoSAR point cloud on the optical image. The bridges are automatically detected from the classification map, by checking the discontinuity of the river followed by morphological operation.

## CONCLUSION

This paper presents the first systematic semantic analysis of meter-resolution spaceborne InSAR point cloud in urban area. Through a strict 3-D fusion of optical images and InSAR point cloud, the attributes extracted from the optical image, e.g. color, semantics, can be rendered to the InSAR point cloud. The complementary information provided by the two data types enables an semantic-level InSAR deformation and 3-D analysis.

The whole algorithm can said to be three steps: reconstruction, fusion, and rendering. The fusion requires both 3-D reconstruction from optical and SAR images, respectively. Optical structure from motion and SAR tomographic inversion are employed for the 3-D reconstruction from the two types of images, respectively. In the fusion step, considering the anisotropic InSAR estimate, an anisotropic iterative closest point algorithm with robustly estimated covariance matrix was introduced. In the rendering part, we introduced a soft point visibility measure which is an improvement of the original hidden point removal algorithm.

## ACKNOWLEDGEMENT

This work was supported by the Helmholtz Association under the framework of the Young Investigators Group “SiPEO” (VH-NG-1018, [www.sipeo.bgu.tum.de](http://www.sipeo.bgu.tum.de)), International Graduate School of Science and Engineering, Technische Universität München (Project 6.08: “4D City”) and the German Aerospace Center (DLR, Förderkennzeichen 50EE1417). The authors would like to thank Dr. Heiko Hirschmüller of DLR-RM for providing the optical data.

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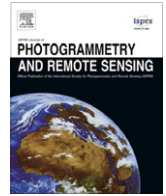


# Appendices

- D Wang Y, Zhu X, Bamler R (2012) Retrieval of Phase History Parameters from Distributed Scatterers in Urban Areas Using Very High Resolution SAR Data. ISPRS Journal of Photogrammetry and Remote Sensing, 73: 89 - 99.**

<http://www.sciencedirect.com/science/article/pii/S0924271612001219>





# Retrieval of phase history parameters from distributed scatterers in urban areas using very high resolution SAR data

Yuanyuan Wang<sup>a,\*</sup>, Xiao Xiang Zhu<sup>a,b</sup>, Richard Bamler<sup>a,b</sup>

<sup>a</sup> Lehrstuhl für Methodik der Fernerkundung, Technische Universität München, Arcisstraße 21, 80333 Munich, Germany

<sup>b</sup> Remote Sensing Technology Institute (IMF), German Aerospace Center (DLR), Oberpfaffenhofen, 82234 Wessling, Germany

## ARTICLE INFO

Article history:  
Available online 1 August 2012

Keywords:  
Phase history  
Distributed scatterer  
Covariance matrix  
SAR  
PSI  
TerraSAR-X

## ABSTRACT

In a recent contribution Ferretti and co-workers (Ferretti, A., Fumagalli, A., Novali, F., Prati, C., Rocca, F., Rucci, A., 2011. A new algorithm for processing interferometric data-stacks: SqueeSAR IEEE Transactions on Geoscience and Remote Sensing 49(9), pp. 3460–3470) have proposed the SqueeSAR method, a way to exploit temporally coherent distributed scatterers in coherent SAR data stacks. Elevation and deformation or subsidence estimates are obtained with accuracy similar as in the well known persistent scatterer interferometry (PSI).

In this paper we propose an alternative approach and provide a first demonstration of the optimal estimation of distributed scatterers' phase histories in urban areas. Different to SqueeSAR, we derive phase histories for each distributed scatterer pixel rather than for groups of pixels. We use the Anderson–Darling statistical test to identify neighboring samples of the same distribution. Prior to covariance matrix estimation required for maximum likelihood estimation we apply a multi-resolution defringe technique. By using TerraSAR-X high resolution spotlight data, it is demonstrated that we are able to retrieve reliable phase histories and motion parameter estimates from distributed scatterers with signal-to-noise-ratio far below the common range.

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## 1. Introduction

Persistent scatterer interferometry (PSI) (Ferretti et al., 2001; Kampes, 2006) exploits the phase history of strong and long term stable scatterers, i.e. the so-called persistent scatterers (PSs), over the entire period of monitoring. Commonly, PSs are selected according to their amplitude dispersion index or their signal-to-clutter ratio (SCR). They are usually seen as bright pixels in synthetic aperture radar (SAR) images. However these criteria are only indications but not necessarily measurements of the decorrelation. Common PSs in real world are stable objects such as window sills on building façades, building corners, and exposed rocks. By proper modeling the phase histories of PSs, the elevation and the time series of the deformation of each PS can be estimated. PSI can provide both large-scale and long-term deformation monitoring with an accuracy of up to mm/year. This is particularly true for urban areas where the PS density is high. As a price for its efficiency, first, PSI considers only a single dominant target within a SAR resolution cell for each PS which forbids layover separation; and second, only permanently reliable scatterers are exploited. This makes PSI an

opportunistic approach and functionally limits the density of the useful scatterers.

The following development of SAR interferometry (InSAR) techniques basically targets at understanding the interaction of the SAR signal within a resolution cell, and at optimally exploring coherent objects other than detected PSs, such as layovered scatterers and distributed scatterers (DSs).

In the first aspect, the authors of (Ferretti et al., 2001) extended PSI by considering phase models for multiple scatterers within a pixel (Ferretti et al., 2005). Or to be more general, differential SAR tomography (D-TomoSAR) (Lombardini, 2005; Fornaro et al., 2009; Zhu and Bamler, 2010, 2011) provides a general solution for such case and avoids the pre-selection of coherent scatterers. D-TomoSAR can separate multiple scatterers at different elevation with possible motion inside a pixel by means of spectral estimation. D-TomoSAR is able to retrieve topography, deformation parameters, or even the entire phase history of multiple scatterers within a resolution cell. Yet compared to PSI, D-TomoSAR is relatively computationally expensive. In the second aspect, the small baseline subset (SBAS) technique (Casu et al., 2005) overcomes the second disadvantage of PSI by further exploiting “temporary” PSs. That is to say, instead of looking for permanently coherent scatterers with respect to a single master acquisition, SBAS takes

\* Corresponding author. Tel.: +49 8153 28 2386.

E-mail address: [yuanyuan.wang@dlr.de](mailto:yuanyuan.wang@dlr.de) (Y. Wang).

advantage of high coherence in pairs of acquisitions with small baselines (both temporal and spatial). In a popular notion, SBAS builds up interferograms between “good” pairs of acquisitions. Then the processing of each set of interferograms with a common master follows standard PSI procedures. Based on these differential results from numbers of interferograms, an inversion of an over-determined linear equation system gives a complete time series of the phase history of a scatterer. SBAS theoretically increases the density of useful scatterers, and decreases the number of images required (Casu et al., 2005).

However, in non-urban environment where surface or volume scattering dominates, PSs density can be very low and DSs are the majority. Even in urban areas, asphalt roads and concrete surfaces seen as DS in X-band also make up a significant percentage. The question then comes to optimally harvest the information from a DS, i.e. how to retrieve its phase history, and model parameters such as elevation and deformation velocity. The challenges are the following. DSs tend to decorrelate temporally, i.e. interferometric DS pixels have usually much higher phase noise than PSs. Also decorrelation may be different for each of the interferograms in the stack.

The SqueeSAR technique (Ferretti et al., 2011a, 2011b) provides a possible solution for these problems. It uses the fact that a DS pixel is usually surrounded by many pixels from the same distribution, also called “brother pixels” in (Ferretti et al., 2011a). SqueeSAR detects these brotherhood segments, uses them for estimation of the covariance matrix and obtains the maximum likelihood estimation (MLE) of the phase history of the segment (De Zan and Rocca, 2005; Seymour and Cumming, 1994). Since SqueeSAR treats each brotherhood segment as a single object, spatial resolution of the estimated phase history is compromised.

This article presents the first demonstration of exploiting DS for urban monitoring using very high resolution SAR image stacks. We estimate elevation and motion history of each DS pixel in an optimum way. Following the basic concept of SqueeSAR, we introduce several new aspects:

- Instead of the brotherhood segments as single objects we retrieve the phase history for each DS pixel.
- For identifying the brotherhood pixels we use a more robust statistical test, i.e. the Anderson–Darlington (AD) test.
- For covariance matrix estimation we introduce a multi-resolution interferometric phase flattening (defringe) algorithm to determine the underlying topographic and motion-induced phase components.

The remainder of this article starts with the explanation of our covariance matrix estimation procedures, followed by phase history retrieval and its model parameters estimation by MLE. Results using TerraSAR-X (TS-X) high resolution spotlight data are present as last.

## 2. Covariance matrix estimation

The covariance between two single-look complex (SLC) measurements  $i_n$  and  $i_k$  in image  $n$  and  $k$  is defined as the expectation of the product of one with the complex conjugate of the other:

$$c_{n,k} = E(i_n i_k^*) \quad (1)$$

Normalizing the covariance by its standard deviation yields the complex correlation coefficients, usually referred to as coherence (Bamler and Hartl, 1998):

$$\gamma_{n,k} = \frac{E(i_n i_k^*)}{E(|i_n|)E(|i_k|)} \quad (2)$$

The MLE of the covariance is obtained by assuming ergodicity and replacing the expectation with averaging  $M$  independent samples:

$$\hat{c}_{n,k} = \frac{1}{M} \sum_m i_n(m) i_k^*(m) \quad (3)$$

and for coherence:

$$\hat{\gamma}_{n,k} = \frac{\sum_m i_n(m) i_k^*(m)}{\sqrt{\sum_m |i_n(m)|^2 \sum_m |i_k(m)|^2}} \quad (4)$$

The covariance matrix  $\mathbf{C}$  containing all the covariances is then

$$\mathbf{C} = \mathbf{Z}\mathbf{Z}^H \quad (5)$$

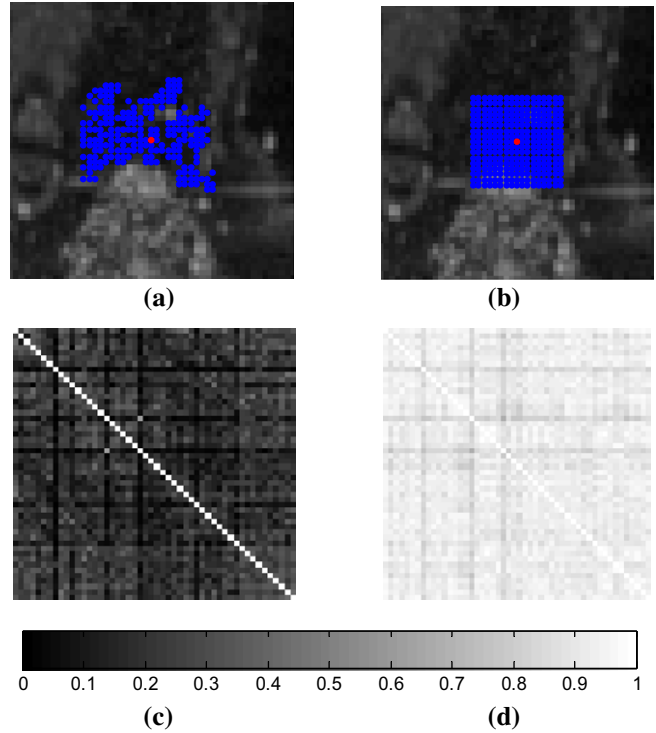
where

$$\mathbf{Z} = \begin{bmatrix} i_1(1) & i_1(2) & \cdots & i_1(M) \\ i_2(1) & i_2(2) & \cdots & i_2(M) \\ \vdots & \vdots & \ddots & \vdots \\ i_N(1) & i_N(2) & \cdots & i_N(M) \end{bmatrix} \quad (6)$$

However, this estimator is biased at low magnitudes of coherence for small  $M$  (Touzi et al., 1999; Bamler and Hartl, 1998). According to the Reed–Mallett–Brennan (RMB) detection loss theory (Reed et al., 1974), to maintain a loss of likelihood (see Eq. (7)) of the estimates of less than 3 dB,  $M$  should be larger than  $2N$ , where  $N$  is the number of measurements:

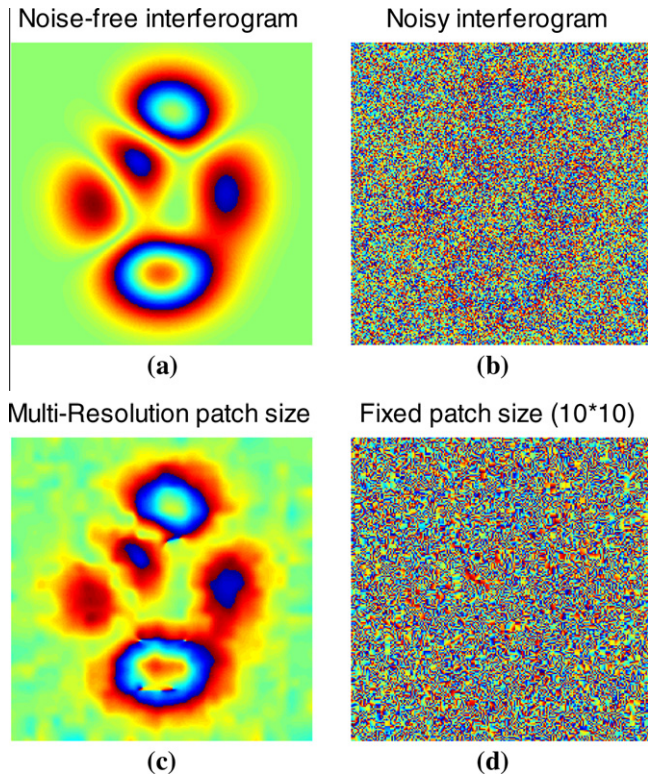
$$\text{loss} = -10 \log_{10}[(M+2-N)/(M+1)] \quad (7)$$

In our experiment, we try to ensure  $M = 3N$  as much as possible. For low coherence bias correction, we adopt a method explained in (Zebker and Chen, 2005) which makes use of an empirical function of estimated coherence with respect to the theoretical one.



**Fig. 1.** (a) and (b): Samples selected by adaptive and rectangle sample selection strategy, respectively. The red dot is the target pixel, while those in blue are the selected pixels, i.e. The neighboring pixels sharing the same statistical distribution as the target pixel. (c) and (d): Corresponding coherence matrices.





**Fig. 2.** (a) Simulated noise-free interferogram, (b) simulated noisy interferogram. 30% Of the pixels are DS with coherence = 0.3 corresponding to a SNR of  $-3.7$  dB in the two SLC images, and the rest 70% are noisy pixels with coherence close to 0, (c) fringes estimated by adaptive multi-resolution method, and (d) fringes estimated using single fixed patch size ( $10 \times 10$  pixels).

Other than this systematic drawback of the covariance estimator, it still faces two major practical issues. Especially in high resolution SAR images of urban areas, the ergodicity of the selected samples is hardly ensured. This jeopardizes the covariance estimation in two ways. First, carelessly selecting these samples, such as using a simple rectangular estimation window (commonly known as box-car method) brings the risk of averaging pixels of different

distributions, and hence different objects, yielding a wrong estimate of the covariance matrix. And second, the spatially varying phase within the estimation window caused by topography or motion diminishes the summation in Eqs. (3) and (4), leading to an underestimation of the covariance or coherence. Therefore, this phase needs to be removed before the covariance estimation. The covariance estimator in Eqn. (3) then becomes:

$$\hat{c}_{n,k} = \frac{1}{M} \sum_m i_n(m) i_k^*(m) \exp(-j(\bar{\phi}_n(m) + \bar{\phi}_k(m))) \quad (8)$$

where  $\bar{\phi}_n$  and  $\bar{\phi}_k$  are the expected phases of  $i_n$  and  $i_k$ .

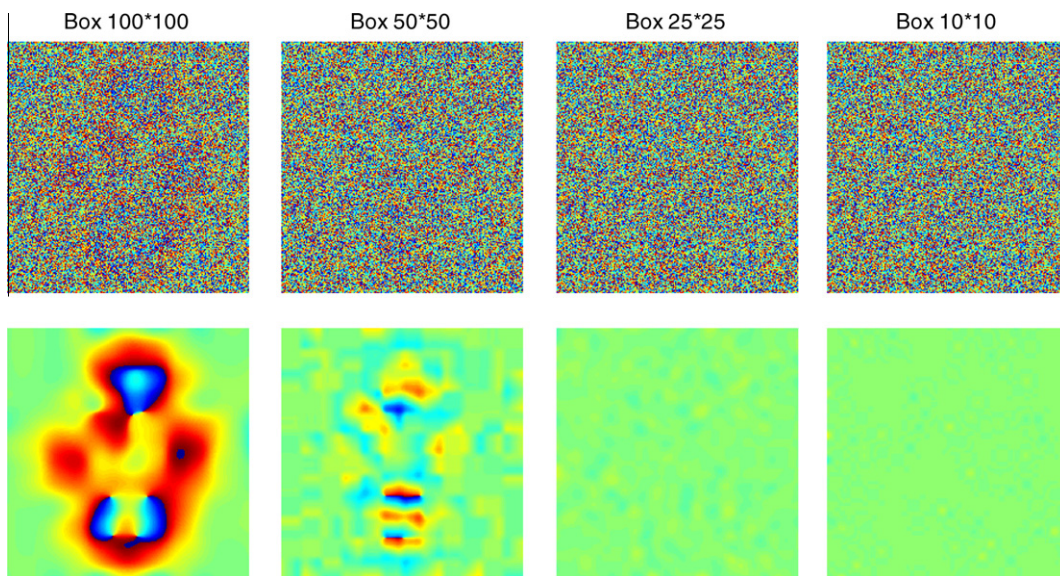
In Eq. (3), the phase of  $c_{n,k}$  is the expected phase difference of  $i_n$  and  $i_k$ . On one hand, we must remove the deterministic part of the phase of  $i_n$  and  $i_k$  for better covariance estimation, and on the other hand we are exploring this interferometric phase from  $c_{n,k}$  for model parameters estimation.

To summarize, the covariance estimation procedure is a two-step approach: (1) adaptive sample selection by AD test on the amplitudes of the samples, and (2) multi-resolution defringe to remove the underlying expectation of the phases. These two steps will be discussed in detail in the following two sections.

### 2.1. Adaptive sample selection

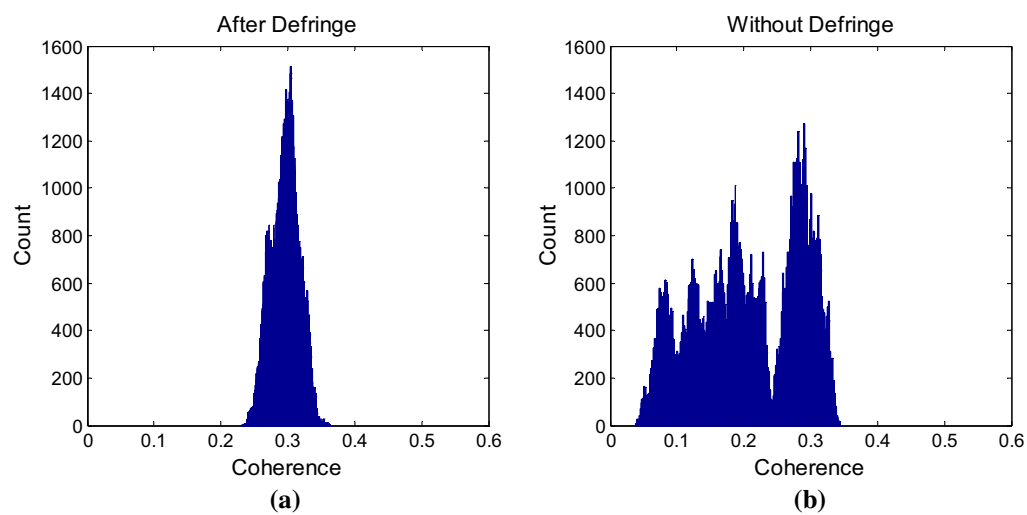
The complex response of a DS in a SAR image is commonly modeled as circular Gaussian variate that widely holds for medium or lower resolution SAR data. However, due to limited number of natural scatterers in a resolution cell in high resolution SAR systems, not fully developed speckle results an imperfect circular Gaussian distribution. Yet we still favor the Gaussian model, as the best asymptotical one, for conveniences in the estimation (i.e. only the covariance is needed).

The MLE of each DS pixel value in the stack is found via inversion of its covariance matrix. As mentioned before, estimation of the covariance matrix elements requires an ensemble of samples of the same distribution. Selecting samples using a rectangular window is justified sometimes, if the rectangle size is small enough and loss of resolution is affordable. However, this is never the case for high resolution images of urban areas, e.g. any PS in the estimation window will dominate the covariance or the coherence estimate of the surrounding pixels. Not only will the high resolution

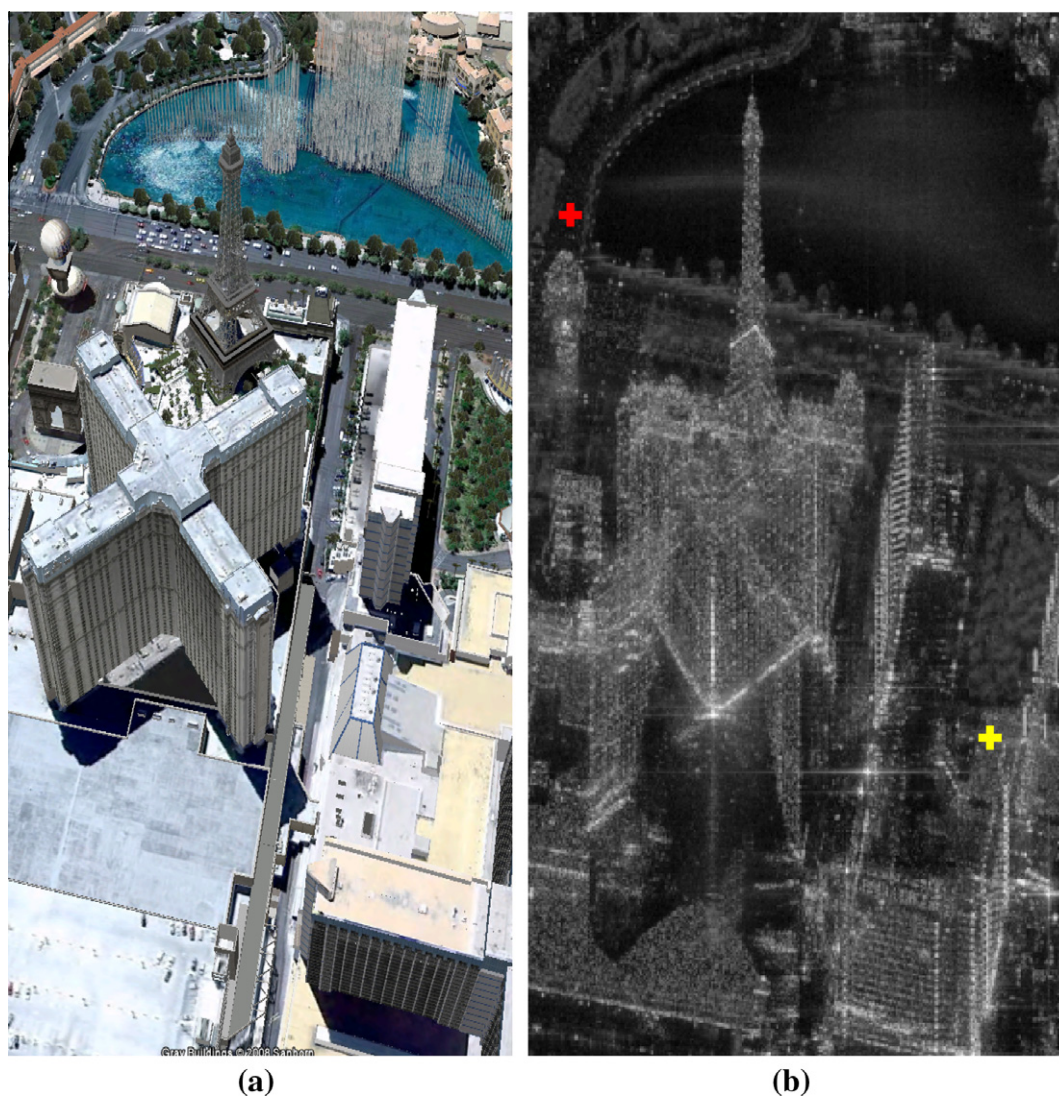


**Fig. 3.** Input noisy interferogram (first row) and estimated fringes (second row) at each resolution level. The estimated fringes are then subtracted from the noisy interferogram. The residual phase is taken as the input for the next resolution level with smaller window size.

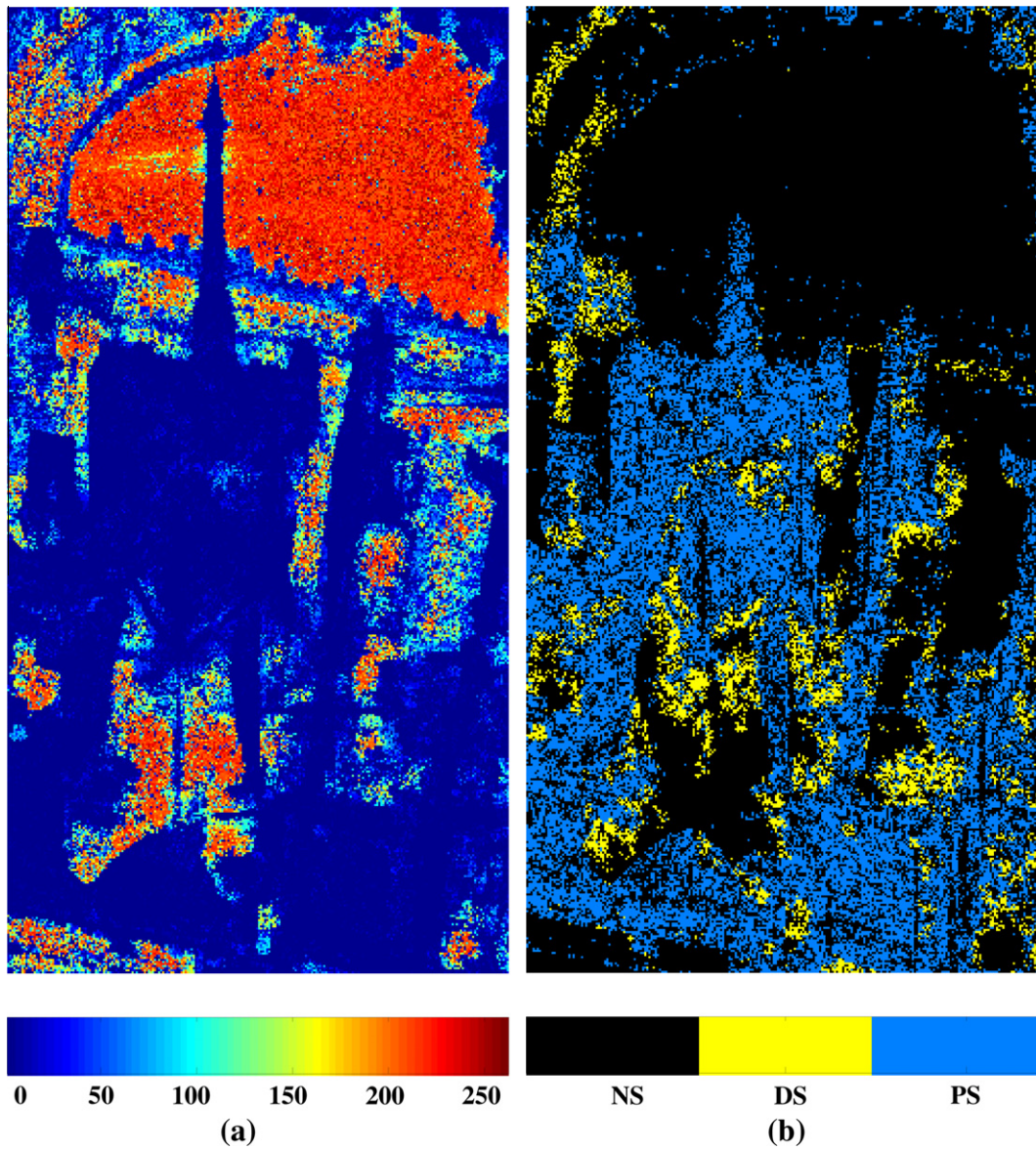




**Fig. 4.** Histograms of coherence estimates of total 107,133 DSs in the simulated interferogram (a) with defringe performed which stick close to the preset value 0.3, and (b) without defringe performed which severely underestimates the coherence.



**Fig. 5.** (a) Optical image with 3D building model from Google Earth™, (b) mean SAR amplitude of the test site. The red and yellow crosses mark the analyzed pixels in Fig. 8.



**Fig. 6.** (a) Number of brother pixels detected within a  $20 \times 20$  pixel box for each pixel of the area shown in Fig. 5. The number is color coded. Red indicates a large number of samples and blue indicates a small number, (b) pixel classification map. The blue pixels indicate the PSs, the yellow the DSs, and the black the NS.

be compromised, but the covariance or coherence of a DS close to a PS will also be severely overestimated.

In (Ferretti et al., 2011a) this problem is tackled by introducing the *DespeckKS* method, which detects the samples adaptively by performing the Kolmogorov–Smirnov (KS) test. The KS test compares the maximum distance between the empirical distribution functions (EDFs) of two data sets which, in our case, are the intensities of one pixel throughout the whole stack with another pixel. By setting a threshold on the distance, the hypothesis that the two pixels belong to the same distribution is accepted or rejected. However, the maximum distance of two EDFs usually happens in the midway of the curve, regardless of the shape of the distribution. Therefore, the KS test is not sensitive to unknown distributions with variations on the tails.

We adopt this idea of amplitude based test. Further considering the fact mentioned in the beginning of this section that fully developed speckle can be compromised in high resolution image and Rayleigh distribution is therefore often not satisfied (Jakeman and Pusey, 1976; Jao, 1984), more sophisticated non-parametric

tests are more appropriate for our purpose. In (Parizzi and Brcic, 2010), different tests are evaluated. Among them, the AD test offers the best detection rate at a constant false alarm rate. Therefore, we replace the KS test by the AD test.

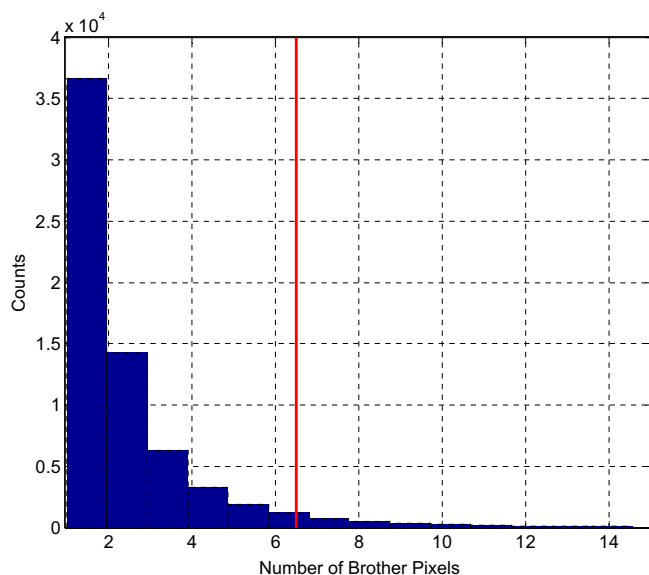
The AD test belongs to the class of quadratic EDF statistics. To compare the probability distributions of pixels  $a$  and  $b$ , we first compute the EDFs  $F_a(x)$  and  $F_b(x)$  of the amplitude values of the pixels through the stack. Then the AD test is a weighted L2-norm of the difference between the two EDFs:

$$A^2 = \frac{N}{2} \sum \frac{[F_a(x) - F_b(x)]^2}{F_{a \cup b}(x)[1 - F_{a \cup b}(x)]} \quad (9)$$

where  $F_{a \cup b}(x)$  is the EDF of the combined pixel values.  $N$  again represents the number of images in our stack. The denominator  $F_{a \cup b}(x)[1 - F_{a \cup b}(x)]$  places more weight on the head and tails of the distributions.

The test is performed for every pixel to identify neighboring pixels of the same distribution, i.e. the brother pixels. These form





**Fig. 7.** Histogram of the number of brother pixels of PSs selected by thresholding the amplitude dispersion index. 95% of the PSs have less than 6 brother pixels. The red line indicates is the cut-off of 95% of all the PSs.

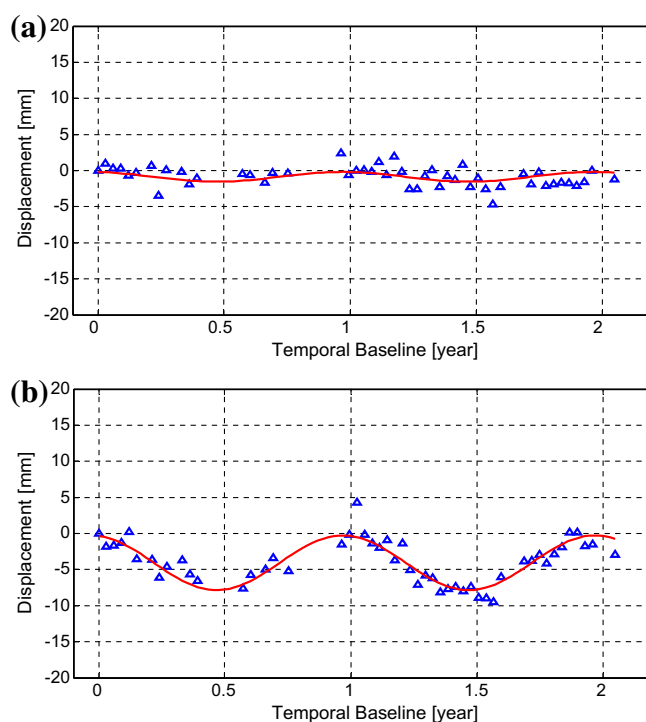
the estimation window. The window size and shape can vary from one pixel to another. This is unlike segmentation where each pixel belongs to a unique class. In our algorithm one pixel is included in different brotherhoods.

Fig. 1a is an example of adaptive sample selection using the AD test on a stack of high resolution TS-X images of downtown Las Vegas. The test area is near the red cross marked in Fig. 5b. The red dots in Fig. 1a and b are our target DS pixel located on the asphalt road between two structures, and the blue ones are the detected brother pixels. As a comparison, samples selected by a rectangular window are marked in Fig. 1b. The adaptive window follows the structure boundary well, while using a rectangle includes all the pixels from both the structures and the road, regardless of their varying amplitudes. The corresponding coherence matrices estimated using the selected samples are shown in the second row of Fig. 1. Since the target pixel is on an asphalt surface having a very low SNR, its coherence is expected to be low. Yet, using the rectangular window, the coherence is heavily overestimated ( $>0.7$ ), as shown in Fig. 1d. Obviously, we are more confident with the results using the adaptive selection window.

## 2.2. Adaptive multi-resolution defringe

Defringe, also known as phase flattening, is to remove an estimate of the expectation, i.e. a low-pass version, of the interferometric phase before covariance estimation (Eq. (8)). In most of the cases a constant local fringe frequency in range and azimuth directions is assumed. The fringe frequency is found by simply searching the maximum coefficient in the Fourier transform of local patches (Bamler and Hartl, 1998; Zebker and Chen, 2005).

Yet the assumption of a constant frequency is not always valid, especially in urban areas with fast varying topography. And the local fringe frequency cannot be correctly estimated at areas where decorrelation noise dominants. We make use an adaptive multi-resolution defringe algorithm, firstly introduced for phase unwrapping (Davidson and Bamler, 1999). It overcomes the aforementioned two problems by on one hand demodulating the interferogram multiple times by estimating local fringes in patches of difference sizes, and on the other hand taking into account only the pixels with sufficiently high coherence.



**Fig. 8.** (a) Phase history of one DS located on the ground, indicated by the red cross in Fig. 5 (b). Its coherence equals 0.16. This pixel experiences no significant deformation, (b) phase history of another DS located on the roof of a building, which undergoes a seasonal motion with 4.5 mm in amplitude. Its coherence is 0.29. This pixel is indicated as the yellow cross in Fig. 5 (b).

We test this algorithm on a simulated interferogram of  $600 \times 600$  pixels. The simulated interferometric phase is shown in Fig. 2b. The pixels in the interferogram are either DS with coherence equal to 0.3, i.e.  $\text{SNR} = -3.7$  dB in both SLC images, or noise (NS) with coherence close to 0. We uniformly distribute 30% of the pixels as DS, and the rest 70% to be NS. This setting simulates the worst real case scenario where no high coherence scatterer is available, but only low coherence targets are sparsely scattered in the scene. The noise-free and noisy interferogram are collocated in Fig. 2a and b. The fringe pattern is hardly distinguishable in the noisy interferogram.

We apply the multi-resolution algorithm to the noisy interferogram. Fig. 3 shows the input and estimated fringes at each resolution level. Starting from the largest patch size, the fringe frequency is estimated using only the DSs in a patch. The interferogram is then demodulated, and passed to the next resolution level. In our test, we set four resolution levels:  $100 \times 100$ ,  $50 \times 50$ ,  $25 \times 25$ , and  $10 \times 10$  pixels. The summation of the phase contributions at all the resolution levels gives the final result in Fig. 2c. For comparison, Fig. 2d shows the result for the usual fixed window size and without employing the pre-selection of coherent pixels. This method completely fails to reconstruct the original interferometric phase, while the multi-resolution algorithm reduces the estimation error tremendously and retains higher correlation with the noise-free interferogram.

After defringing, the coherences of all the DSs in the interferogram are evaluated. The results show consistent and accurate solutions with respect to the preset coherence of 0.3. 99% of the results lie in the interval of  $0.3 \pm 0.05$ . This result is compared to the coherence estimated without defringe. Fig. 4 plots the histogram of the estimated coherence, with defringe and without defringe, of total 107,133 DSs. It is obvious that the coherence is severely underestimated if defringe is not performed.

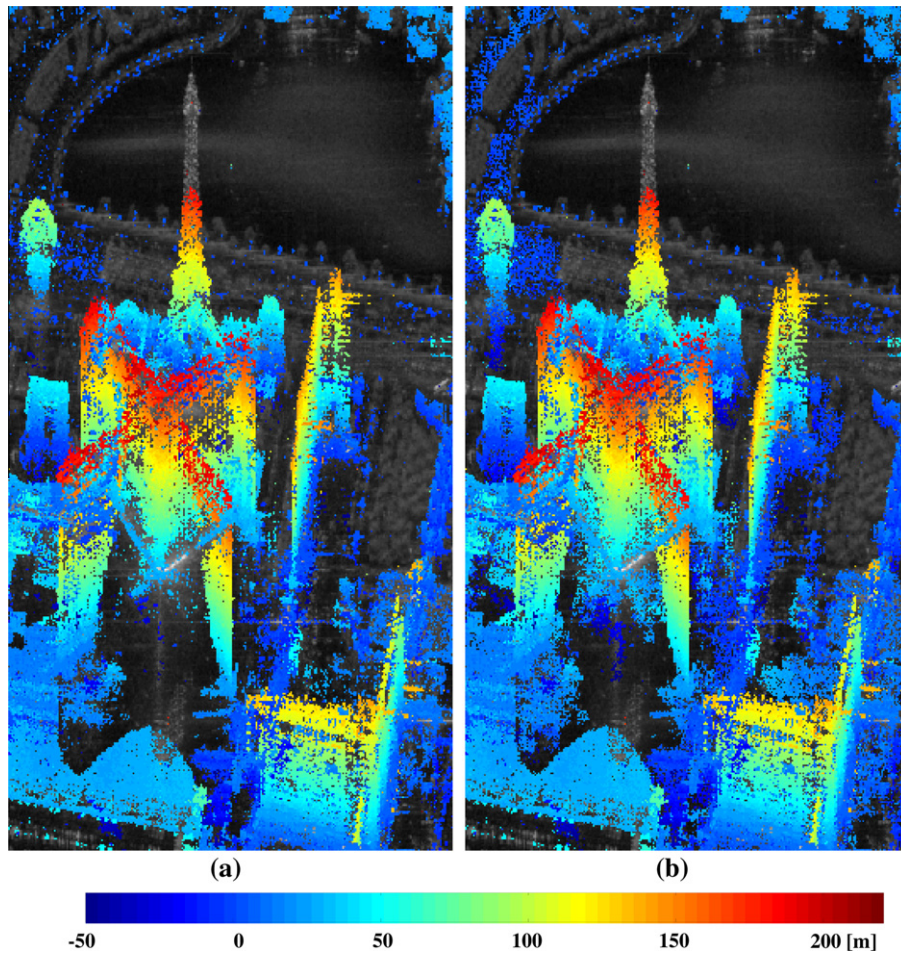


Fig. 9. Estimated elevation at (a) only PSs, and (b) both PSs and DSs.

### 3. Parameters estimation

#### 3.1. Maximum likelihood estimator

We use MLE for parameter estimation for each DS pixel. We make use of the objective function developed in (De Zan and Rocca, 2005):

$$\{\hat{h}, \hat{a}\} = \arg \max \left\{ \frac{\exp(-\mathbf{i}^H \Phi(h, a) \mathbf{C}_0^{-1} \Phi(h, a)^H \mathbf{i})}{\det[\mathbf{C}_0] \pi^N} \right\} \quad (10)$$

where  $\mathbf{i}$  is the complex measurements vector;  $\Phi$  is diagonal matrix containing the modeled phase;  $\mathbf{C}_0$  is absolute value of the covariance matrix;  $h, a$  is height above reference surface, line-of-sight (LOS) motion parameter such as linear deformation velocity and seasonal displacement amplitude and  $N$  is the number of images

In the likelihood function, the covariance matrix  $\mathbf{C}$  is decomposed to  $\Phi(h, a) \mathbf{C}_0^{-1} \Phi(h, a)^H$ . The diagonal matrix  $\Phi$  contains the phase of the model. The denominator  $\det[\mathbf{C}_0] \pi^N$  can be ignored: it is a constant for each pixel.

### 4. Experiments

#### 4.1. Test site 1 – Paris Las Vegas hotel

We apply the proposed algorithm to a test site about  $500 \times 500$  m centered at Paris Las Vegas hotel in downtown Las Vegas cropped from a stack of co-registered 50 TS-X high resolution spotlight images. Fig. 5a and b are the optical image from Google

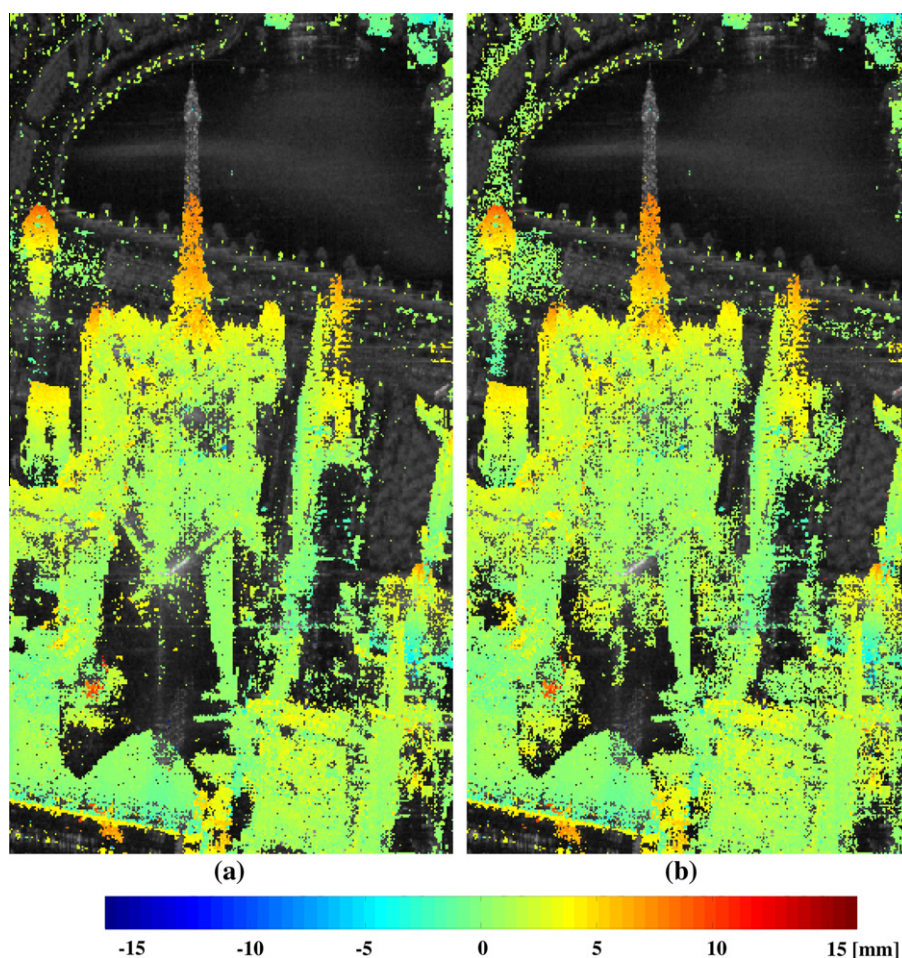
Earth™ and the mean amplitude image of the test area, respectively. In this test case, the area consists of different types of scattering objects, i.e. partly building façades (probable PS candidates), partly asphalt roads and rough surface (probable DS candidates), and a large fraction of totally decorrelated scatterers (e.g. the fountain on the top of the scene, and vegetated areas).

The pre-processing, including amplitude calibration and atmospheric phase screen (APS) correction, is performed using the PSI-GENESIS system from the German Aerospace Center (DLR) (Kampes, 2006; Adam et al., 2003). We firstly classify all the pixels to PS, DS, or NS, by jointly considering the coherence and the number of brother pixels. Based on the classification, we process PSs using the standard PSI technique, and DSs using the proposed algorithm.

#### 4.2. Pixel classification

To classify PS, DS and NS, we firstly identify the PSs by making use of the property explained in (Ferretti et al., 2011a) that most of the PSs possess very few brother pixels. Showing in Fig. 6, the number of the brother pixel map, most of the building structures show up in blue which indicates a small number of brother pixels, in contrast to the orange color for the rest of the image. Statistics tell that 95% of the PSs (selected by thresholding on amplitude dispersion index) have less than 6 samples (see Fig. 7). Secondly, the DSs are distinguished from the rest of the pixels by thresholding on the coherence. The threshold is set at the unbiased lower limit of the coherence estimator in Eq. (4) using 200 looks. The final classifica-





**Fig. 10.** Estimated seasonal deformation amplitude at (a) only PSs, and (b) both PSs and DSs.

tion is shown in Fig. 6b, where PS, DS, and NS are marked as different colors. In this test site, the number of DSs is about 20% of PSs.

#### 4.3. Retrieved phase history

According to the classification map, we estimate the covariance matrix for each DS after sorting all the interferograms along the temporal baseline. The phases of the covariance matrix's first column represent the expected interferometric phases referring to the first image. Since defringe was applied prior to the covariance estimation, the pre-removed phase was added back to get the final phase history of each pixel. Fig. 8 plots the phase histories of two DSs (indicated as red and yellow crosses in Fig. 5b). Seen from that, the first one is located on the asphalt road at ground level, with presumably no significant deformation. The reflectivity of such material is generally very low, as we can see the surrounding pixels are consistently dark. For this pixel, the coherence is 0.16, equivalent to an average SNR of  $-7.1$  dB. Its phase history shows almost no deformation throughout the two years' time span. The second one is on the roof of a building, possibly subject to thermal expansion. It has a better coherence of 0.29, corresponding to an average SNR of  $-4$  dB. It undergoes a seasonal deformation of 4.5 mm in amplitude. The red curves in the figures are the fitting of the sinusoidal deformation model.

#### 4.4. Estimated parameters

Following the retrieved phase history, we estimate the elevation and seasonal motion amplitude for each DS. The result is combined with the estimates from PSs using the standard PSI

procedures. All the estimates are referred to a single reference point. Figs. 9a and 10a give the estimates from PSs only, and Figs. 9b and 10b are PSs and DSs combined. We see a linear gradient from the base of the buildings to the top in Fig. 9, with blue indicating low elevation, and red for high. A similar gradient can also be observed in the seasonal motion amplitude estimate in Fig. 10. The seasonal motion amplitude becomes more positive as the elevation gets higher representing displacement towards the sensor. This is mostly caused by the natural expansion and contraction of the building itself, but also subject to the effect of height dependent APS to a maximum of about 1.5 mm per 100 m in vertical height (Cong and Eineder, 2012).

The PSs already provide very good coverage over most parts of the building façades, except for the upper half of the tower due to its very complex motion. The DSs give extra information over the rest of the image. The contributions of DSs are scattered over the whole area, including areas on the upper left corner and area close to the center of the image. Most of the DSs are on the ground level, indicated by the blue color. Their deformation amplitudes are generally small (green). It is worthy to point out that seemingly reliable estimates also found in the shadow area. They are “ghost scatterers” caused by mirror reflections of the façade at the ground (Auer et al., 2011). The variances of these DS estimates are comparable with those of PS estimates, although the coherences of these DSs are much lower. Nevertheless, the MLE of the phase history of DS is obtained by covariance matrix estimation based on many samples. The residual of the phase history to the sinusoidal model is about 1–2 mm, thus providing a reliable estimation of the model parameters.



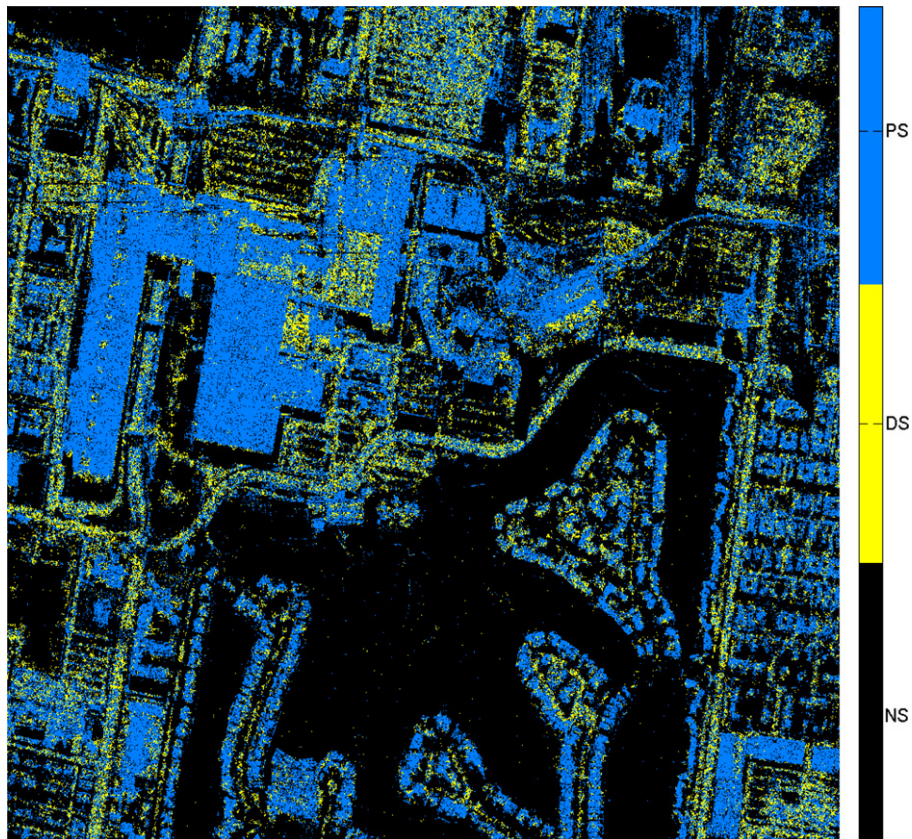


Fig. 11. Pixel classification map. The blue pixels indicate the PSs, the yellow the DSs, and the black the NS.

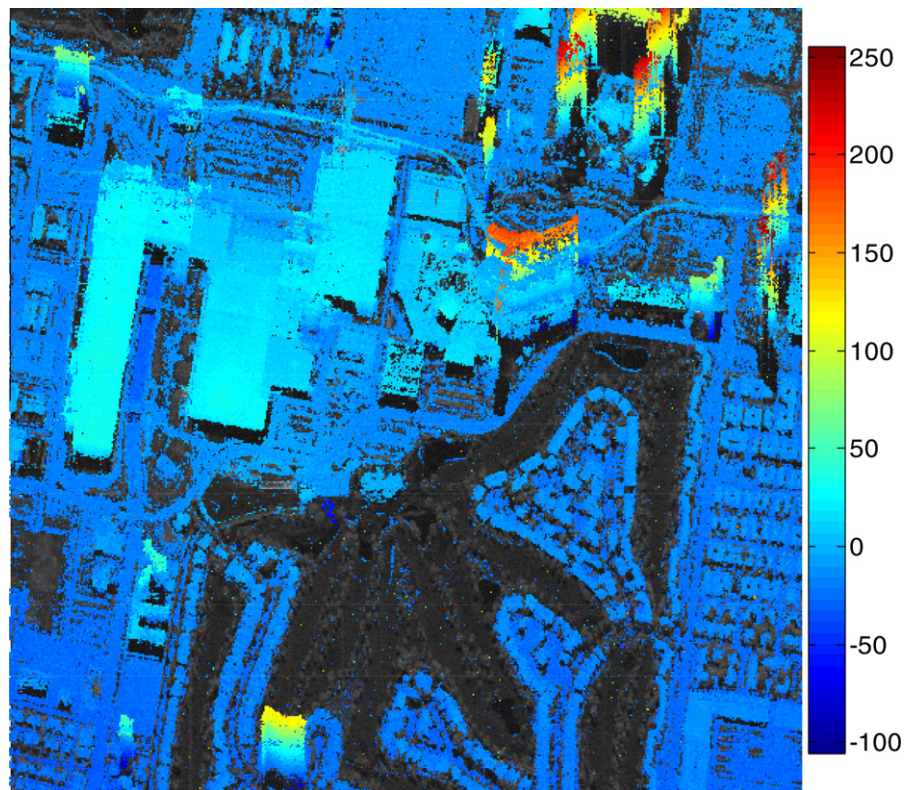


Fig. 12. Estimated elevation of both PSs and DSs of the second test site. The unit is in meter.



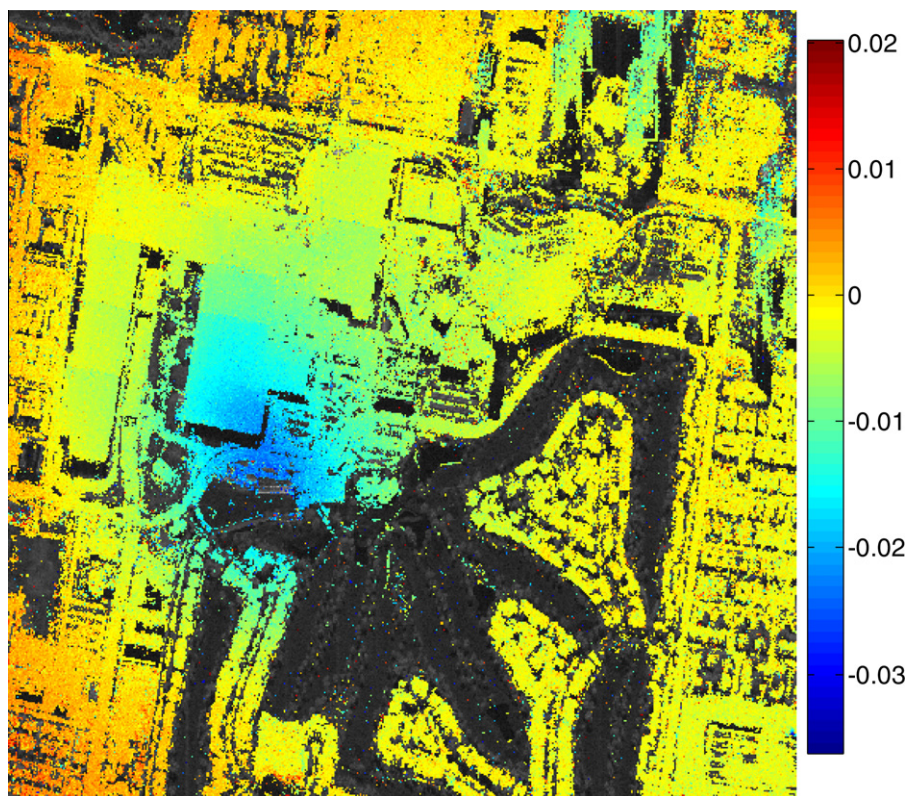


Fig. 13. Estimated linear deformation rate of both PSs and DSs. The unit is in m/year.

#### 4.5. Test site 2 – Las Vegas convention center

A larger area about  $1 \times 2$  km around the Las Vegas convention center is chosen as the second test site due to a known subsidence discovered in (Zhu and Bamler, 2011). The same processing procedures are applied. The pixel classification is shown in Fig. 11. Blue color (PS) fills up most of the building structures, while yellow (DS) appears mostly on the road. The park located on the lower half of the image is almost black due to temporal decorrelation of vegetation. In Figs. 12 and 13, the estimation results of PSs and DSs combined are demonstrated. The elevation estimates look consistent and reliable for both PS and DS, except part of the building façades due to the layover problem. The linear deformation rate estimates show a subsidence pattern centered at Las Vegas convention center with a maximum of about 15 mm/year.

#### 5. Conclusion and outlook

This article presents optimal DS phase history and model parameters retrieval in urban areas using very high resolution TS-X spotlight data. To preserve the high spatial resolution, particular focus is put on the phase history retrieval for each DS pixel. We emphasize the critical problem of accurate covariance matrix estimation. A solution is provided by using adaptive sample selection and a multi-resolution defringe method. The adaptive sample selection method is proven to be an essential procedure in the covariance matrix estimation. The AD test performs well in distinguishing targets of different scattering characteristics. The adaptive multi-resolution defringe algorithm outperforms the method with a fixed window size in terms of fringe estimation accuracy and robustness to noise. The effectiveness of the proposed method is tested using simulated and real TS-X data. Periodic seasonal motion histories of sample DS have been derived. The residuals to the sinusoidal model were in the order of 1–2 mm.

In the future, as a more general solution, we will improve the DS covariance matrix estimation using a “fuzzy brotherhood” concept, i.e. using surrounding pixels with weightings derived from a statistical test instead of using a fixed threshold. The improved algorithm will be applied to non-urban area, and possible mixed single- and repeat-pass data stacks.

#### Acknowledgements

This work is supported by International Graduate School of Science and Engineering (IGSSE), Technische Universität München (IGSSE Project 6.08: “4D City”), and the German Research Foundation (DFG, Förderkennzeichen BA2033/3-1). The authors would like to thank Alessandro Parizzi from DLR for valuable discussions and one of the reviewers who pointed out the possible misinterpretation of height dependent APS as seasonal motion.

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# Appendices

- E Wang Y, Zhu X (2015c) Robust Estimators in Multipass SAR Interferometry. IEEE Transactions on Geoscience and Remote Sensing, 54 (2): 1 - 13.**

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# Robust Estimators for Multipass SAR Interferometry

Yuanyuan Wang, *Student Member, IEEE*, and Xiao Xiang Zhu, *Senior Member, IEEE*

**Abstract**—This paper introduces a framework for robust parameter estimation in multipass interferometric synthetic aperture radar (InSAR), such as persistent scatterer interferometry, SAR tomography, small baseline subset, and SqueeSAR. These techniques involve estimation of phase history parameters with or without covariance matrix estimation. Typically, their optimal estimators are derived on the assumption of stationary complex Gaussian-distributed observations. However, their statistical robustness has not been addressed with respect to observations with nonergodic and non-Gaussian multivariate distributions. The proposed robust InSAR optimization (RIO) framework answers two fundamental questions in multipass InSAR: 1) how to optimally treat images with a large phase error, e.g., due to unmolded motion phase, uncompensated atmospheric phase, etc.; and 2) how to estimate the covariance matrix of a non-Gaussian complex InSAR multivariate, particularly those with nonstationary phase signals. For the former question, RIO employs a robust M-estimator to effectively downweight these images; and for the latter, we propose a new method, i.e., the *rank M-estimator*, which is robust against non-Gaussian distribution. Furthermore, it can work without the assumption of sample stationarity, which is a topic that has not previously been addressed. We demonstrate the advantages of the proposed framework for data with large phase error and heavily tailed distribution, by comparing it with state-of-the-art estimators for persistent and distributed scatterers. Substantial improvement can be achieved in terms of the variance of estimates. The proposed framework can be easily extended to other multipass InSAR techniques, particularly to those where covariance matrix estimation is vital.

**Index Terms**—Differential interferometric synthetic aperture radar (D-InSAR), SAR interferometry (InSAR), *M*-estimator, rank covariance matrix, robust InSAR optimization (RIO), robust estimation.

## I. INTRODUCTION

### A. State-of-the-Art Multipass InSAR Techniques

MULTIPASS interferometric synthetic aperture radar (InSAR) techniques exploit a time series signal of scatterers. For spaceborne SAR data, they are commonly known as differential InSAR (D-InSAR) as multiple acquisitions are

usually acquired by repeat passes. Well-known techniques include persistent scatterer interferometry (PSI) [1]–[3], differential SAR tomography [4]–[6], small baseline subset [7], [8], SqueeSAR [9], [10], CAESAR [11], and so on. They aim at retrieving the parameters of interest, such as the scatterer's 3-D position and deformation rate, by utilizing persistent scatterers (PSs) and/or distributed scatterers (DSs). They are widely employed for ground deformation monitoring, as well as individual building and urban 3-D infrastructure reconstruction. In general, millimeter accuracy of the yearly linear deformation rate and meter accuracy of the scatterer's 3-D position can be achieved with respect to a reference point [1], [12], [13].

However, these accuracy levels refer to the optimal estimators derived based on the assumption of Gaussian-distributed data. PS is modeled as a deterministic signal with additive zero-mean white complex circular Gaussian (CCG) noise [1], whereas DS is modeled as correlated zero-mean CCG [9], [14], [15]. Deviation from the assumption will greatly compromise the performance of the estimator. It boils down to two fundamental problems in the parameter estimation of multipass InSAR.

- Unmodeled phase such as residual atmospheric/orbit phase errors, unmodeled motion phase, etc: Such phase errors impose non-Gaussian noise on PS and DS.
- Non-Gaussian scatterers that bias the covariance matrix estimation of DS: Their spatial stationarity cannot be guaranteed because of the spatially varying phase.

For the first type, there has not been a systematic method to optimally weight images with large phase error. The performance of current techniques depends on the modeling of the interferometric phase, which includes the deformation model, the preprocessing step of atmospheric/orbit error phase mitigation, and so on. This is true for any multipass InSAR processor that targets large area processing. The common deformation models are usually linear and periodic models. The atmospheric phase mitigation methods are mainly based on PS and its variants [1], [16]–[18]. Their accuracy depends on the density of the network, the number of images, and the validity of the assumed deformation model. For meter-resolution data, particularly in the urban areas, the atmospheric phase can be usually well mitigated. Other methods are available based on weather data [19]–[21], which can overcome some disadvantages of the PS-based method, e.g., the topography-correlated atmospheric phase. In [20] and [22], it was concluded that centimeter accuracy can be achieved for controlled locations on the ground. However, achieving centimeter accuracy for a network of dense points on a city scale for InSAR deformation monitoring purposes is not yet feasible at present. Only until recently that such concept and preliminary result have been presented as the *Geodetic TomoSAR* [23].

For the second type, the prevailing approach is to make amplitude-based adaptive sample selections using statistical

Manuscript received April 16, 2015; revised July 20, 2015; accepted August 14, 2015. Date of publication September 14, 2015; date of current version January 19, 2016. This work was supported in part by the Helmholtz Association under the framework of the Young Investigators Group “SiPEO” under Grant VH-NG-1018, by the International Graduate School of Science and Engineering, Technische Universität München under Project 6.08: “4-D City,” and by the German Aerospace Center (DLR) under Förderkennzeichen 50EE1417. (Corresponding author: Xiao Xiang Zhu.)

Y. Wang was with the International Graduate School of Science and Engineering, 85748 Garching, Germany. He is now with the Signal Processing in Earth Observation, Technische Universität München, 80333 Munich, Germany (e-mail: wang@bv.tum.de).

X. X. Zhu is with The Remote Sensing Technology Institute, German Aerospace Center (DLR), 82234 Weßling, Germany, and also with the Signal Processing in Earth Observation, Technische Universität München, 80333 Munich, Germany (e-mail: xiao.zhu@dlr.de).

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Digital Object Identifier 10.1109/TGRS.2015.2471303

tests such as Kolmogorov–Smirnov (KS) or Anderson–Darling tests [8]–[10], [24]. However, estimation results with these methods degrade as the number of images decrease. According to [25], assuming a Rayleigh-distributed amplitude time series of *ten* observations, the detection rate is only 20% (at a constant 5% false alarm rate) for distinguishing two DSs with an expected intensity ratio of 3 dB using the KS test. As detection rate decreases, pixels of different distributions, including those non-Gaussian, are included in the covariance matrix estimation. This in turn affects the final parameter estimation.

### B. RIO Framework

The aim of this paper is to introduce robust estimation for multipass InSAR techniques when the underlying statistical model of the observations is not fulfilled, which was briefly mentioned in [26] and [27]. Let  $f_{\mathbf{g}}(\mathbf{g}|\boldsymbol{\theta}, F(\hat{\mathbf{C}}))$  be a generic likelihood function of the observation time series  $\mathbf{g}$ , the parameter vector  $\boldsymbol{\theta}$  to be retrieved, and the system model  $F(\cdot)$  (with the estimated covariance matrix  $\hat{\mathbf{C}}$ , if used). The maximum-likelihood estimator (MLE) is  $\hat{\boldsymbol{\theta}} = \arg \max_{\boldsymbol{\theta}} f_{\mathbf{g}}(\mathbf{g}|\boldsymbol{\theta}, F(\hat{\mathbf{C}}))$ . Assuming a CCG-distributed  $\mathbf{g}$ , the MLE can be simplified to the minimization of the sum of the squared residuals, i.e.,

$$\hat{\boldsymbol{\theta}} = \arg \max_{\boldsymbol{\theta}} \ln f_{\mathbf{g}}(\mathbf{g}|\boldsymbol{\theta}, F(\hat{\mathbf{C}})) = \arg \min_{\boldsymbol{\theta}} \|\boldsymbol{\varepsilon}\|_2^2 \quad (1)$$

where  $\boldsymbol{\varepsilon}$  is the residual vector, and  $\|\cdot\|_2$  denotes the Euclidean norm.

The proposed robust method introduces the following aspects into the current techniques.

- It replaces the MLE with an *M*-estimator [28], which minimizes the sum of a customized function  $\rho(x)$  of the residuals, i.e.,

$$\hat{\boldsymbol{\theta}} = \arg \min_{\boldsymbol{\theta}} \sum_i \rho(\varepsilon_i). \quad (2)$$

- Should DS be used, the covariance matrix estimator is also replaced with its robustified version. Depending on the stationarity of the samples' phases, different estimators are proposed for:
- Stationary samples: *M*-estimator of covariance that is an amplitude-based weighting of the sample covariance matrix;
- Nonstationary samples: *rank M*-estimator (RME) that is an *M*-estimator using the *rank* of the complex observations  $\mathbf{g}$ .

Table I summarizes a comparison of the current state-of-the-art approaches with the improvements proposed by the RIO framework.

### C. Readers' Guide

In this paper, Section II covers the necessary basics of the *M*-estimator and the multivariate rank. Sections III and IV introduce and provide detailed information about the robust estimators used in InSAR techniques, followed by experiments using simulated and real data in Section V.

Notations of some frequently used variables and their dimensions are listed in the following. Italic small letters denote scalar

TABLE I  
COMPARISON OF RIO FRAMEWORK WITH THE CURRENT APPROACH

		State-of-the-art	RIO
Covariance estimation			
DS	Stationary	Sample covariance	M-estimator: $\hat{\mathbf{C}}_{M-est}$
	Nonstationary	matrix: $\hat{\mathbf{C}}_{MLE}$	Rank M-estimator: $\hat{\mathbf{C}}_{RME}$
Phase-history parameters estimation			
PS		Periodogram: $\hat{\boldsymbol{\theta}}_{MLE}$	M-estimator: $\hat{\boldsymbol{\theta}}_{M-est}$
DS	Stationary	Periodogram on samples mean: $\hat{\boldsymbol{\theta}}_{MLE}$	M-estimator on samples mean: $\hat{\boldsymbol{\theta}}_{M-est}(\hat{\mathbf{C}}_{M-est})$
	Nonstationary	–	M-estimator pixel-wise inversion: $\hat{\boldsymbol{\theta}}_{M-est}(\hat{\mathbf{C}}_{RME})$

value, bold small letters are vectors, and bold capital letters denote a matrix.

$N \in \mathbb{N}$	number of observations;
$M \in \mathbb{N}$	number of samples;
$\mathbf{g} \in \mathbb{C}^N$	complex InSAR observations;
$\mathbf{r} \in \mathbb{C}^N$	rank of the observations;
$\boldsymbol{\theta} \in \mathbb{R}^L$	<i>L</i> parameters of the phase model;
$\mathbf{C} \in \mathbb{C}^{N \times N}$	complex covariance matrix;
$\boldsymbol{\Phi} \in \mathbb{C}^{N \times N}$	diagonal matrix of the modeled phase;
$\mathbf{W} \in \mathbb{R}^{N \times N}$	diagonal weighting matrix;
$\boldsymbol{\varepsilon} \in \mathbb{C}^N$	residuals;
$\rho(x)$	loss function;
$ \cdot $	on scalar: absolute value; on vector and matrix: element-wise absolute value
$\ \cdot\ $	vector norm;
$\det(\cdot)$	matrix determinant.

## II. BACKGROUND INFORMATION

### A. *M*-Estimator Basics

*M*-estimators are a class of well-known robust estimators that take the form of a minimization problem:  $\min \sum_i \rho(x_i)$ . The *M*-estimator permits a minimization of a customized *loss function*  $\rho(x)$  of the residuals to resist outliers without preprocessing the data, e.g., by outlier trimming.

The derivation of the *M*-estimator can be described as follows. Consider a simple location estimation problem, where the likelihood function of the univariate observation of the location  $g \in \mathbb{R}$  is  $f_g(g|\mu)$  in which  $\mu$  is the expected location to be estimated.  $f_g(g|\mu)$  equals a probability density function (pdf)  $f_x(x)$ , where  $x = g - \mu$ . Without losing generality, let us assume that  $x$  is scaled to unit variance. Maximizing the sum of log-likelihood is the MLE of  $\mu$ , given multiple observations of  $g$ , i.e.,

$$\hat{\mu}_{MLE} = \arg \min_i -\ln f(x_i). \quad (3)$$

The  $M$ -estimator of  $\mu$  is as follows:

$$\hat{\mu}_{M\text{-est}} = \arg \min_i \sum \rho(x_i). \quad (4)$$

Therefore, choosing  $\rho(x) = -\ln f_x(x)$  yields the ordinary MLE. For example, the MLE, when assuming a normal distribution, corresponds to an  $M$ -estimator with  $\rho(x) = x^2$ .

Solving (4) is equivalent to finding the zero position of its derivative, which can be written as

$$\sum_i \frac{\rho'(x_i)}{x_i} x_i = 0 \quad (5)$$

where  $\rho'(x)$  is the derivative of  $\rho(x)$ .  $\rho'(x)/x$  is commonly referred to as the weighting function  $w(x)$  because substituting  $x_i = g_i - \mu$  into the previous equation yields an  $M$ -estimator that is actually a weighted average, i.e.,

$$\hat{\mu}_{M\text{-est}} = \frac{\sum_i w(g_i - \hat{\mu}) g_i}{\sum_i w(g_i - \hat{\mu})}. \quad (6)$$

The weighting function of a given pdf  $f_x(x)$  can be derived with the following relationship:

$$w(x) = \frac{-\partial \ln f_x(x)}{x \partial x}. \quad (7)$$

Equation (6) is solved iteratively with an initial  $\mu$  estimate being given and the weights being updated at each iteration. Typically, the weighting function is designed to have high efficiency at normal distribution. Therefore, if  $x$  is not scaled to unit variance, the weights are  $w = \rho'(x/\sigma)/(x/\sigma)$ , where  $\sigma$  is the standard deviation of  $x$ . Convergence is only guaranteed for convex  $\rho$  functions [29]. The initial estimate must be carefully estimated should  $\rho$  be nonconvex.

The  $M$ -estimator is a tradeoff between robustness and efficiency, where the efficiency of an estimator refers to the closeness of the variance of its estimates to the possible lower bound, e.g., the Cramér–Rao lower bound for an unbiased estimator [30]. Nevertheless, it can still maintain high efficiency under the nominal distribution by properly choosing the loss function.

### B. Multivariate Rank

Another important tool in robust estimation is the *rank* of random variables. Using the ranks of the samples instead of the samples themselves reveals their robustness when assessing the statistical dependence between two random variables, and more importantly, it is invariance to the nonlinear relation between the two random variables. For example, the Spearman's rank correlation coefficient is the ordinary correlation coefficient of the samples' ranks. It is robust against nonstationary samples, i.e., the nonlinear transformation between two random variables [31].

In univariate real random variables, the rank simply refers to the integer ranking of each realization of this variable according to its value. In [32], the centered rank of the sample  $x_i$  in the univariate data set  $\{x_i, x_2, \dots, x_M\}$  is defined as

$\hat{r}_i = (1/M) \sum_{j=1}^M \text{sign}(x_i - x_j)$ , where  $\text{sign}(x) = -1, 0, 1$  for  $x < 0$ ,  $x = 0$ , and  $x > 0$ , respectively. This is simply a ranking algorithm that can be easily extended to real multivariate case, i.e.,

$$\hat{\mathbf{r}}_i = \frac{1}{M} \sum_{j=1}^M \text{sign}(\mathbf{x}_i - \mathbf{x}_j) = \frac{1}{M} \sum_{j=1}^M \frac{\mathbf{x}_i - \mathbf{x}_j}{\|\mathbf{x}_i - \mathbf{x}_j\|} \quad (8)$$

for the sample set  $\{\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_M\}$ , where  $\mathbf{x}_i \in \mathbb{R}^N$  is a vector of observations, such as a time series of pixels in multi-pass InSAR techniques, despite the fact that it is not a complex number in this case.

## III. ROBUST COVARIANCE ESTIMATION

The covariance estimate of a target pixel requires a selected neighborhood of samples. The widely used sample covariance matrix is only suitable for samples following a stationary CCG distribution. Otherwise, a robust estimator is necessary. In the following, we demonstrate the robust covariance estimators for both stationary and nonstationary samples.

### A. Sample Covariance Matrix

If a complex SAR pixel value  $g$  obeys the classic model of a zero-mean CCG distribution [14], then multiple acquisitions  $\mathbf{g} \in \mathbb{C}^N$  at the location of the same pixel lead to a multivariate CCG distribution, i.e.,

$$f_{\mathbf{g}}(\mathbf{g}) = \frac{1}{\pi^N \det(\mathbf{C})} \exp(-\mathbf{g}^H \mathbf{C}^{-1} \mathbf{g}) \quad (9)$$

which is fully characterized by its covariance matrix  $\mathbf{C}$ , where  $N$  is the number of acquisitions. Given a set of  $M$  spatially stationary samples  $\mathbf{G} = [\mathbf{g}_1, \mathbf{g}_2, \dots, \mathbf{g}_M]$ , the MLE of  $\mathbf{C}$  is one that maximizes the likelihood, i.e.,

$$\hat{\mathbf{C}}_{\text{MLE}} = \arg \min_{\mathbf{C}} \sum_{m=1}^M \mathbf{g}_m^H \mathbf{C}^{-1} \mathbf{g}_m. \quad (10)$$

This yields the well-known equation for the sample covariance matrix:

$$\hat{\mathbf{C}}_{\text{MLE}} = \frac{1}{M} \sum_{m=1}^M \mathbf{g}_m \mathbf{g}_m^H. \quad (11)$$

$\hat{\mathbf{C}}_{\text{MLE}}$  is not robust against any type of outlier. A single outlier in the samples will considerably bias the estimates. Hereafter, we designate the MLE as the MLE for CCG, unless explicitly stated otherwise.

### B. Robust Estimation for Stationary Non-Gaussian Samples

If the selected samples are non-Gaussian, caused by either a non-Gaussian scattering mechanism or a poor sample selection, the covariance estimation can be robustified using an  $M$ -estimator. Here, we assume the samples' stationarity.

The  $M$ -estimator of a covariance matrix is expressed analogously with (4) by its extension to multivariate variables, i.e.,

$$\hat{\mathbf{C}}_{M\text{-est}} = \arg \min_{\mathbf{C}} \sum_{m=1}^M \rho(x_m(\mathbf{C})) \quad (12)$$

where the squared residual  $x_m^2(\mathbf{C}) \in \mathbb{R}$  of the multivariate  $\mathbf{g}_m$  is  $\mathbf{g}_m^H \mathbf{C}^{-1} \mathbf{g}_m$ . It can be solved using an iteratively reweighted approach [33], [34], i.e.,

$$\hat{\mathbf{C}}_{k+1} = \frac{1}{M} \sum_{m=1}^M w(x_m(\hat{\mathbf{C}}_k)) \mathbf{g}_m \mathbf{g}_m^H. \quad (13)$$

The real-valued weighting function  $w(x)$  is  $\rho'(x)/x$ , and  $k$  denotes the iteration index. The weighting function downweights highly deviating samples whose whitened version  $x_m(\hat{\mathbf{C}}_k) = \|\hat{\mathbf{C}}_k^{-1/2} \mathbf{g}_m\|$  is large in magnitude, which greatly depends on the intensities of  $\mathbf{g}_m$ .

To derive a meaningful weighting function for some heavily tailed distributions, we model the non-Gaussian SAR pixel distribution as a complex  $t$ -distribution [35], [36], i.e.,

$$f_{\mathbf{g}}(\mathbf{g}) = \frac{\Gamma(\nu + N)}{(\pi\nu)^N \Gamma(\nu) \det(\mathbf{C})^N} \left(1 + \frac{1}{\nu} \mathbf{g}^H \mathbf{C}^{-1} \mathbf{g}\right)^{-(\nu+N)} \quad (14)$$

where  $\Gamma(\cdot)$  is the gamma function, and  $\nu \in \mathbb{R}^+$  is the degree of freedom (DoF). The complex  $t$ -distribution is a realistic model for SAR observations as the DoF allows a quantitative definition of its deviation from CCG. This distribution approaches CCG as the DoF approaches  $+\infty$  and becomes more heavily tailed as it decreases toward zero. In practical terms, this provides flexibility for handling different levels of outliers.

The weighting function can be derived according to (7). However, (14) must be transformed into its equivalent real-valued version [33] to perform the derivation. Therefore, we form a new multivariate  $\tilde{\mathbf{g}} = [\text{Re}(\mathbf{g}); \text{Im}(\mathbf{g})]$  ( $\text{Re}[\cdot]$  and  $\text{Im}[\cdot]$  are the real and imaginary parts of a complex number; semicolon denotes vertical concatenation), which follows a  $2N$ -variate real-valued  $t$ -distribution [33]. The corresponding weighting function can be derived thereafter as follows (see also Appendix A):

$$w(x) = \frac{2N + \nu}{\nu + 2x^2}. \quad (15)$$

For good overall robustness, a small  $\nu (< 5)$  is preferred [37]. For example, in [38]  $\nu$  is chosen from 2.5 to 3.9 for centimeter-resolution millimeter-wave SAR data of urban areas. The particular complex  $t$ -distribution considering  $\nu = 1$  is also known as the complex Cauchy distribution, which has been thoroughly investigated in many studies.

To best safeguard against unknown heavily tailed distributions,  $\nu$  can be set to zero. This leads to a weighting function of  $w(x) = N/(\mathbf{g}_m^H \mathbf{C}^{-1} \mathbf{g}_m)$ . Further, by assuming  $\mathbf{C}$  equals  $\bar{I} \mathbf{I}$ , where  $\bar{I}$  is the expected intensity, and  $\mathbf{I}$  is the identity matrix, the iterative process can be dropped. This leads to the interesting weighting function  $w = \bar{I} \|\mathbf{g}_m\|^{-2}$ . The corresponding covariance matrix estimate

$$\hat{\mathbf{C}}_{\text{SCM}} = \frac{N\bar{I}}{M} \sum_{m=1}^M \|\mathbf{g}_m\|^{-2} \mathbf{g}_m \mathbf{g}_m^H \quad (16)$$

is also known as the *sign covariance matrix* (SCM) [32], [39], where only the direction of each multivariate sample is considered. The real covariance is lost as a result. Should the

exact covariance be retrieved, one could always estimate it from the eigenvectors of the SCM and the samples. SCM does not require iteration and is robust against outliers.

SCM is an engineering solution for fast processing under the general  $M$ -estimator's framework. There also exist other methods such as amplitude clipping and dropping, etc. The choice among the methods depends on the data and application. For DSs with low coherence, simulation shows SCM outperforms amplitude clipping and dropping both in terms of variance and bias.

### C. Robust Estimation for Nonstationary Non-Gaussian Samples

To complete the theory, we must consider the situation in which samples are neither stationary nor complex Gaussian. The most common nonstationarity found in an SAR interferogram is the uncompensated rapidly varying interferometric phase among the spatial samples. For nonstationary samples, a joint estimation of the expected interferometric phase and the covariance [15] is required, i.e.,

$$\hat{\mathbf{C}}_{\text{MLE}} = \frac{1}{M} \sum_{m=1}^M \hat{\Phi}_m^H \mathbf{g}_m \mathbf{g}_m^H \hat{\Phi}_m \quad (17)$$

where  $\hat{\Phi}_m$  is the complex diagonal matrix containing the estimated phase values of  $\mathbf{g}_m$ .

Usually, the estimation of interferometric phase and the covariance matrix is separated. The former is usually achieved by common band filtering by either spatial averaging or spectral trimming. Its performance greatly depends on the filter itself; additionally, estimations can be challenging in urban areas. In [10], a multiresolution defringe algorithm to manage this challenge is used. In any case, the phase estimation requires additional effort, and poor estimates significantly degrade the covariance matrix estimation.

This calls for an estimator that is invariant for multiplicative phase signals. As demonstrated in [32] and [39], the rank covariance matrix for real multivariate is invariant with regard to the additive deterministic signal. Inspired by this fact, we introduce a new estimator, i.e., RME, for complex multivariate with a multiplicative phase signal. By replacing the multivariate  $\mathbf{g}$  with its rank  $\mathbf{r}$  in (13), we obtain the following:

$$\hat{\mathbf{C}}_{\text{RME},k+1} = \frac{1}{M} \sum_{m=1}^M w(x_m(\hat{\mathbf{C}}_{\text{RME},k})) \hat{\mathbf{r}}_m \hat{\mathbf{r}}_m^H. \quad (18)$$

We define the complex rank vector, being analogous to (8), as follows:

$$\hat{\mathbf{r}}_m = \frac{1}{J} \sum_{j=1}^J \frac{\mathbf{g}_m \cdot \mathbf{g}_j^*}{\|\mathbf{g}_m \cdot \mathbf{g}_j^*\|} \quad (19)$$

where  $\mathbf{g}_j$  is the direct neighborhood of  $\mathbf{g}_m$ , and  $\cdot$  denotes the element-wise multiplication. Note its difference from the original definition, where subtraction is replaced by multiplication of the complex conjugate. The deterministic phase has been reduced through this operation, and it approaches zero through



averaging. To maintain effective phase cancelation, the direct neighborhood should be kept small. We usually restrict it to only four adjacent pixels. The RME is a fourth-order descriptor of the sample statistics. It can be proven (Appendix B) that the elementwise square root of  $|\hat{\mathbf{C}}_{\text{RME}}|$  approaches  $|\hat{\mathbf{C}}_{\text{MLE}}|$  asymptotically under a CCG distribution when using a one-neighborhood sample. The variance of the square root  $|\hat{\mathbf{C}}_{\text{RME}}|$  is also higher than that of  $|\hat{\mathbf{C}}_{\text{MLE}}|$  because of the multiplication of two uncorrelated complex-valued random variables. Therefore, the number of samples should be doubled when using RME. Hereafter, we assume that the elementwise square root on  $|\hat{\mathbf{C}}_{\text{RME}}|$  has been performed.

#### IV. RETRIEVAL OF ROBUST PHASE HISTORY PARAMETERS

##### A. PS Phase History Parameters

One of the commonly used estimators for PS phase history parameters  $\theta$  is the periodogram [40] (often with amplitude-dropped measurements [1], [3]), i.e.,

$$\hat{\theta} = \arg \max_{\theta} \left\{ \left| \frac{1}{N} \sum_{n=1}^N g_n \exp(-j\varphi_n(\theta)) \right| \right\} \quad (20)$$

where  $g_n$  and  $\varphi_n$  are the complex pixel value and the modeled phase of the  $n$ th image, respectively. Usually, the amplitude of  $g_n$  is dropped in the estimation, as it barely changes estimates at high SNR. Equation (20) is the MLE when assuming additive independent and identically distributed (i.i.d.) CCG noise [41], i.e.,

$$\hat{\theta} = \arg \max_{\theta} \left\{ (2\pi\sigma^2)^{-N} \exp \left( -\frac{1}{2\sigma^2} (\mathbf{g} - \bar{\mathbf{g}}(\theta))^H (\mathbf{g} - \bar{\mathbf{g}}(\theta)) \right) \right\} \quad (21)$$

where  $\sigma$  and  $\bar{\mathbf{g}}(\theta)$  are the standard deviation of the CCG distribution and the modeled PS pixel values, respectively.

Based on the aforementioned MLE, we propose the following robust estimator of PS phase history parameters to deal with possible large phase errors, particularly the residual atmospheric phase and unmodeled motion phase, i.e.,

$$\hat{\theta} = \arg \min_{\theta} \sum_{i=1}^N \rho \left( \frac{\text{Re}[\varepsilon_i(\theta)]}{\sigma_R} \right) + \rho \left( \frac{\text{Im}[\varepsilon_i(\theta)]}{\sigma_I} \right) \quad (22)$$

where the residual  $\varepsilon_i(\theta)$  is  $g_i - \bar{g}_i(\theta)$ ; and  $\sigma_R$  and  $\sigma_I$  are the standard deviations of real and imaginary parts, respectively. The most appropriate  $\rho(x)$  function should be chosen according to the pdf of the PS noise, which depends on the performance of the atmospheric phase correction, the choice of motion model, etc. If such information is not available, a good alternative is Tukey's biweight loss function, i.e.,

$$\rho(x) = \begin{cases} -\frac{(c^2 - x^2)^3}{6c^4} + \frac{c^2}{6} & |x| < c \\ \frac{c^2}{6} & \text{elsewhere} \end{cases} \quad (23)$$

where  $c \in \mathbb{R}^+$  is a tuning parameter beyond which the loss is constant. The lower the value of  $c$ , the more robust but less

efficient is the estimator, and *vice versa*.  $c$  is usually chosen to be 4.586, which is said to give an efficiency of 95% at normal distribution [29]. For an overview of different robust loss functions, readers are referred to [29] and [42].

Equation (22) should be solved iteratively, where  $\sigma_R$  and  $\sigma_I$  are updated at each iteration. Their initial values are obtained from the initial estimate  $\hat{\theta}_0$ ; this is not critical for convex  $\rho(x)$ . However, a robust initial estimation is strongly recommended for nonconvex  $\rho(x)$ , such as the Tukey's loss function. Fortunately, the initial estimation need not be efficient. To give one example, the least trimmed square that minimizes the sum of  $q$  smallest squared residuals [43] is as follows:

$$\hat{\theta}_0 = \arg \min_{\theta} \sum_{i=1}^q |\varepsilon_i(\theta)|^2. \quad (24)$$

The initial estimates of standard deviation, as well as those at each iteration, can be obtained using a robust estimator such as the median absolute deviation, i.e.,

$$\hat{\sigma} = 1.483 \cdot \text{median}(|\varepsilon - \text{median}(\varepsilon)|). \quad (25)$$

The factor 1.483 is a normalization constant, which gives an expectation of unity for a normal distribution. With this combination, convergence can be achieved in just a few iterations; therefore, the computational cost is approximately three to four times more than that for the standard estimator. The same procedures can be easily extended to the LAMBDA estimator [44] for PS phase parameter estimations, which has a substantial number of users.

##### B. DS Phase History Parameters

Because of the same artifacts caused by the phase calibration mentioned earlier, the estimator of DS phase history parameters must also be robustified. If stationarity is assumed for a DS and its neighborhood, one can treat DSs as a single PS by averaging them, as proposed in SqueeSAR, and then the robustified estimator is identical to (22).

However, we do not assume stationarity in our considerations. Our objective is a full inversion of individual single-look DS observation (DS pixels without averaging). The DS MLE introduced in [15] is recalled here as follows:

$$\hat{\theta} = \arg \min_{\theta} \left\{ \mathbf{g}^H \Phi(\theta) |\hat{\mathbf{C}}|^{-1} \Phi(\theta)^H \mathbf{g} \right\} \quad (26)$$

where  $\Phi(\theta) \in \mathbb{C}^{N \times N}$  is the diagonal matrix containing the modeled phase of  $\mathbf{g}$ . Seemingly, the robustified estimator for DS phase history parameters takes an expression similar to (22), but with the residuals  $\varepsilon(\theta)$  whitened by a robust covariance matrix estimate, e.g.,  $\hat{\mathbf{C}}_{\text{RME}}$ , i.e.,

$$\varepsilon(\theta) = |\hat{\mathbf{C}}_{\text{RME}}|^{-0.5} \Phi(\theta)^H \mathbf{g}. \quad (27)$$

The robustified DS estimator can be written into an iteratively reweighted expression with  $k$  being the iteration index, i.e.,

$$\hat{\theta}_{k+1} = \arg \min_{\theta} \left\{ \varepsilon^H(\theta) \mathbf{W}_k \varepsilon(\theta) \right\} \quad (28)$$

TABLE II  
PSEUDOCODE OF THE ROBUST ESTIMATOR FOR SINGLE-PIXEL DS  
OBSERVATIONS

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Get initial estimate for  $\hat{\boldsymbol{\theta}}$  which can be the same for all DS samples in the same
neighborhood

Calculate initial  $\mathbf{W}_0$ 

for iteration  $k = 1$  until convergence
  for DS sample  $m = 1$  to  $M$ 
    Estimate  $\hat{\boldsymbol{\theta}}^m$  by solving (28) using  $\mathbf{W}_k$ 
  end
  Compute the robust mean residuals  $\bar{\boldsymbol{\varepsilon}}$  of the neighborhood following (29).
  Update  $\mathbf{W}_k$  based on the mean residual
end

```

---

where  $\mathbf{W} \in \mathbb{R}^{N \times N}$  is the diagonal weighting matrix whose element is calculated using each element of  $\boldsymbol{\varepsilon}(\boldsymbol{\theta})$  based on  $w_i = \rho'(\varepsilon_i)/\varepsilon_i$ .

However, we should be aware that the marginal distribution of the phase of zero-mean CCG multivariate is uniform as it is in its whitened version. The atmospheric phase, presumably also uniformly distributed over time, applies no change to the DS observations statistically. In other words, the robust loss function is blind to phase contamination from single-look multivariate DS observations. Therefore, correct weighting of the contaminated observations must be introduced to the estimator. It should be determined by the expected residuals  $\bar{\boldsymbol{\varepsilon}}$  of the neighborhood ensemble. The expected residuals can be calculated by simple averaging. However, because of possible outliers in the neighborhood, it should be estimated robustly as follows:

$$\bar{\boldsymbol{\varepsilon}} = \frac{\sum_{m=1}^M w^m \boldsymbol{\varepsilon}^m(\hat{\boldsymbol{\theta}}^m)}{\sum_{m=1}^M w^m} \quad (29)$$

where the superscript  $m$  denotes the sample number in the neighborhood, and  $w^m$  is a robust weight.

To summarize, (28) is a joint estimation of the phase parameters of individual single-look DS observations in a neighborhood. Its computation should begin with initial estimates of each sample in the neighborhood (assumed to be the same), which jointly determine the initial weighting matrix. This matrix is used to retrieve the parameters of each single-look DS observation time series in the neighborhood and is updated based on all the estimates upon finishing one iteration. The pseudocode of the proposed DS estimator is as stated in Table II.

One practical issue when dealing with real data is the positiveness of the covariance matrix. The inverse square root operation requires that the matrix be positive definite, i.e., for any nonzero vector  $\mathbf{y} \in \mathbb{R}^N$  that  $\mathbf{y}^T |\hat{\mathbf{C}}| \mathbf{y} > 0$ . This criterion is not fulfilled when the rank of the selected samples is less than  $N$ , i.e., when they are spatially correlated, which results in some eigenvalues of  $|\hat{\mathbf{C}}|$  being numerically close to zero. On rare occasions, the covariance matrix must be regularized. In our implementation, this is performed by adaptively adding an

TABLE III  
PSEUDOCODE OF THE COVARIANCE MATRIX REGULARIZATION

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 $\mathbf{I}$  = identity matrix

 $e \in \mathbb{R}^+$  is a small number, e.g., 1e-6

while  $|\hat{\mathbf{C}}_{RME}|$  is NOT positive definite
   $|\hat{\mathbf{C}}_{RME}| = |\hat{\mathbf{C}}_{RME}| + e\mathbf{I}$ 
   $e = e \times 2$ 
end

```

---

appropriately small quantity to its diagonal. The pseudocode is as stated in Table III.

## V. ROBUSTNESS ANALYSIS

### A. Comparison of Robust Covariance Estimators

We compared the original MLE, the  $M$ -estimator with  $t$ -distribution weighting, and the RME under three different scenarios: 1) multivariate CCG; 2) multivariate complex  $t$ -distribution with one DoF; and 3) nonstationary multivariate  $t$ -distribution with one DoF. The data are simulated to have ten acquisitions, with a neighborhood of 1000 samples. Each sample vector is simulated by dewhiting a ten-variate vector of zero-mean i.i.d. CCG distribution (for scenario 1) or complex  $t$ -distribution (for scenarios 2 and 3) with a predefined coherence matrix. In the last scenario, we added linear fringes with ten different fringe frequencies randomly picked within  $[0 \pi/10]$  to the phases of the ten acquisitions.

We also predefined two different coherence matrices: one exponentially decaying and the other with a constant 0.5 coherence between acquisitions. The results are shown in Fig. 1, where the left subfigure corresponds to the exponential coherence matrix, and the right corresponds to the constant matrix. Each row represents one of the aforementioned three scenarios, and each column represents one of the three covariance estimators. The subplots (1, 1) in the two subfigures are regarded as the reference coherence matrices because the MLE is the optimal estimator under CCG and is asymptotically unbiased.

All three estimators can preserve the correct shape of the covariance matrix under CCG. The MLE fails in the second scenario, where the samples are contaminated by outliers. The coherence is usually overestimated because of the large amplitude of the outliers. The  $M$ -estimator and RME remain as correct estimations. In particular, this  $M$ -estimator is the MLE for the simulated data in the second scenario. Finally, the  $M$ -estimator is not capable of dealing with nonstationary samples. Heavy underestimation occurs because of the summation of the complex numbers with nonstationary phases. The RME is mean invariant and maintains good performance in all conditions. However,  $\hat{\mathbf{C}}_{RME}$  experiences higher bias at low coherence [as observed in (a)] and higher variance [as observed in (b)], due to the doubling of the phase variance of the rank observation. Doubling the number of looks should be used in the estimation of  $\hat{\mathbf{C}}_{RME}$ . This has been mentioned in Section III-C. The

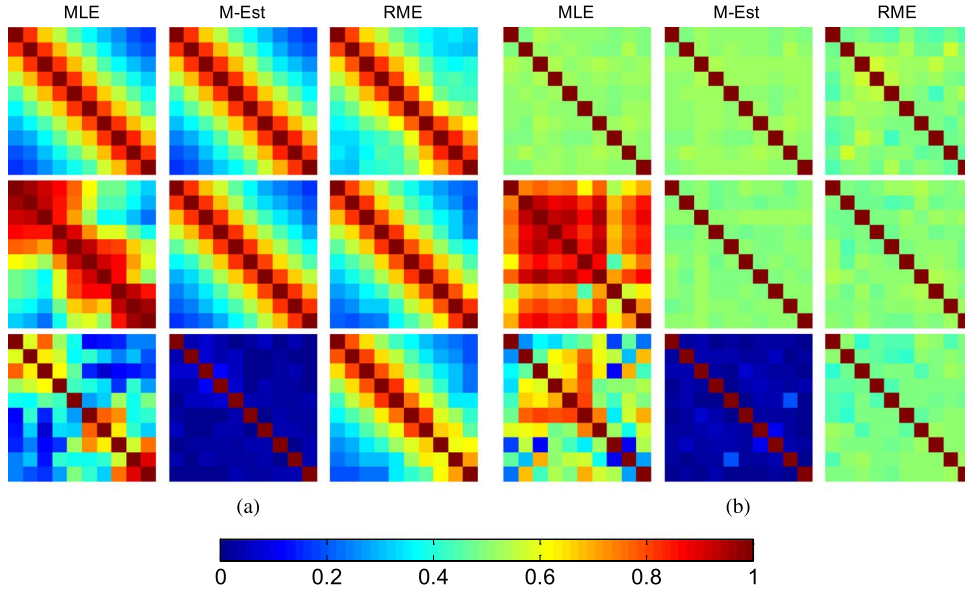


Fig. 1. Comparison of three covariance matrix estimators under three different observation cases. First row: CCG. Second row: Complex  $t$ -distribution with one DoF. Third row: Nonstationary complex  $t$ -distribution with one DoF. (a) and (b) show the results of the exponential and constant coherence matrices, respectively. In both (a) and (b), first column: MLE (under Gaussian); second column:  $M$ -estimator with  $t$ -distribution weighting; and third column: RME with  $t$ -distribution weighting.

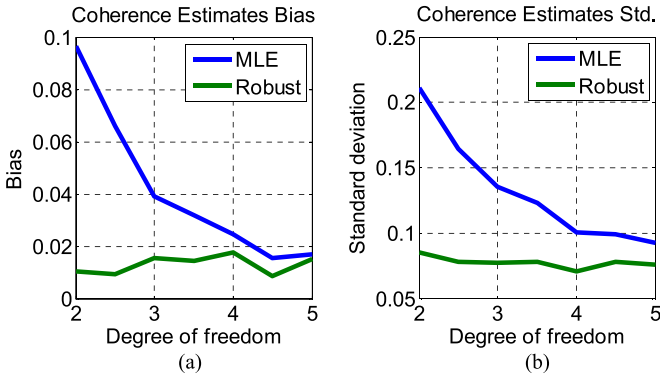


Fig. 2. (a) Bias and (b) standard deviation of the coherence estimates of MLE and the proposed robust estimator for true coherence = 0.2. One hundred looks are used in the coherence estimation.

efficiency of the  $M$ -estimator and that of RME are always lower than 100% at CCG distribution. This illustrates the tradeoff between robustness and efficiency.

To quantitatively analyze the performance of the robust estimator at different outlier levels, we have performed experiments using less tailed distribution by changing the DoF in the complex  $t$ -distribution. The complex  $t$ -distribution approaches complex Gaussian as the DoF increases. Fig. 2 plots the variance and bias of the robust estimator and MLE as a function of the DoF for true coherence = 0.2. As one can see, the robust estimator outperforms the Gaussian MLE. However, as we expected, the advantage of robust estimator becomes less as the DoF increases because the distribution becomes more Gaussian.

In practice, the number of samples should be sufficiently large to prevent any significant bias in the covariance estimates and to reduce the variance of the estimates. We found that 200 samples are more than sufficient for a stack of 30 images

based on the Reed–Mallett–Brennan detection rule [45], which measures the loss of the signal's log-likelihood. When using RME, the number of samples should be doubled to achieve a comparable variance with that of the MLE.

#### B. Efficiencies of Robust PS and DS Estimators

Next, we used a Monte Carlo simulation to quantitatively compare the efficiencies of the proposed PS and DS estimators with their original MLEs. We set the common parameters for the simulation to be similar to those of the TerraSAR-X data, where the wavelength  $\lambda = 0.031$  m, the range distance  $R = 700$  km, the acquisitions number  $N = 20$ , and the spatial baselines  $\mathbf{b}_\perp$  were randomly sampled from  $-100$  m to  $+100$  m with uniform distribution, and the temporal baselines  $\mathbf{t}$  were evenly spaced from  $-1$  year to  $+1$  year.

The PS observation vector is simulated by adding zero-mean i.i.d. CCG noise with a certain SNR to the deterministic PS signal  $\mathbf{g} = A \exp(-j4\pi/\lambda(s\mathbf{b}_\perp/R + vt))$ , where  $A$  is a complex amplitude. The same SNR was maintained for each acquisition. A DS vector was simulated using the same procedure described earlier, i.e., by whitening a multivariate zero-mean i.i.d. CCG using a predefined coherence matrix. The coherence was maintained constant between any two acquisitions. To complete the DS simulation, we added the interferometric phase  $-j4\pi/\lambda(s\mathbf{b}_\perp/R + vt)$  to the phases of the DS vector. We generated 1000 samples of this DS vector for the covariance matrix estimation. These represent the contamination-free PS and DS observations.

We then introduced contamination in two ways. First, we added a random phase to eight (40%) of the 20 acquisitions. The phase was uniformly distributed between  $[-\pi, \pi]$  for both PS and DS. For DS, we added the same phase to the samples of the same acquisition to simulate the temporally random but

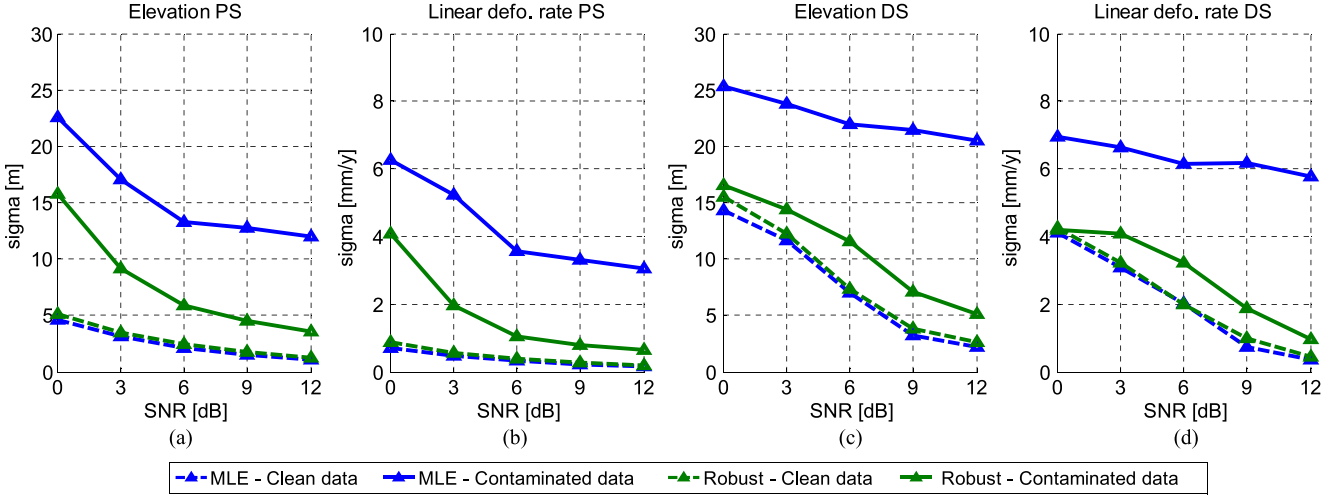


Fig. 3. Comparison of estimates of the standard deviation of elevation and linear deformation rate by the robust estimators and the ordinary MLEs. (a) and (b) refer to the results of the PS estimators, and (c) and (d) refer to the DS estimators. In each subfigure, the green curves are the results of the proposed robust estimators, and the blue curves are those of the original MLE; the dashed curves are the estimators performing on observations with a CCG distribution, and the solid curves are the results for contaminated data, which we generated by adding a uniformly distributed random phase to eight of the 20 acquisitions. The blue dashed curves in all the subfigures are the MLEs under a CCG distribution; hence, they are the optimal estimates.

spatially stationary atmospheric phase. Second, we used an i.i.d. complex  $t$ -distribution with one DoF in the DS simulation to simulate outliers in the DS samples.

Both contamination-free and contaminated observations were used in the proposed robust estimators and their original counterparts. We used the Tukey's loss function as the robust kernel for both robust estimators of PS and DS. Parameter  $c$  was set to 4.586, and we repeated the Monte Carlo simulation 1000 times. Fig. 3 shows the standard deviation of the estimates of elevation and linear deformation rate with respect to different SNRs. Fig. 3(a) and (b) correspond to the PS estimators, and Fig. 3(c) and (d) correspond to the DS estimators. The SNR of the DS refers to the correlated signal w.r.t. the decorrelated part and can be directly related to the coherence using the formula  $\gamma = (1 + \text{SNR}^{-1})^{-1}$  [14], [46]. In each subfigure, the green curves are the results of the proposed robust estimators, and the blue curves are the original MLEs; the dashed curves are the estimators performing on observations with CCG distribution noise, and the solid curves are contaminated data results. The blue dashed curves in all the subfigures are the MLEs under CCG; hence, they are the optimal estimates. All the simulation results suggest that the robust estimators significantly outperform the original MLEs when up to 40% of the observations are contaminated. The robust estimators improve the efficiency by a factor of 7–35 for contaminated data (the squared ratio of the solid curves). Still, the efficiency of the robust estimator at the nominal distribution is close to that of the original MLE. The *relative efficiency* of the robust estimator and the MLE, defined as

$$\eta_{\text{RE}} = \frac{\sigma_{\text{MLE}}^2}{\sigma_{\text{Robust}}^2} \quad (30)$$

at the nominal distribution of the MLE, is approximately 70% for the robust PS estimator and 80% for the DS estimator (the squared ratio of the dashed curves). Therefore, the robust

estimators are also capable of handling uncontaminated observations without too much loss of efficiency.

### C. Real Data Performance

We selected two test sites from a stack of 33 TS-X high-resolution spotlight interferograms of the super volcano Campi Flegrei, Italy. The acquisition time spans from 2009 to 2012. The coregistration and atmospheric phase correction were performed by the German Aerospace Center (DLR)'s *IWAP* processor, i.e., the integrated wide-area processor [47] based on its predecessor *PSI-GENESIS* [2]. Independent experiments were conducted by comparing the linear deformation estimates for evaluating the performance of the robust covariance estimator, the performance of the robust loss function against the uncompensated phase error, and the overall performance over a large area.

1) *Robustness of the RME of Covariance*: Fig. 4 shows the difference between the linear deformation rate of the DSs estimated using the classical sample covariance matrix  $\hat{\mathbf{C}}_{\text{MLE}}$  (left column) and that using the proposed  $\hat{\mathbf{C}}_{\text{RME}}$  (right column). Identical samples were used for estimating these two covariance matrices. They were adaptively selected with the KS test using *ten* amplitude images, which we expected to introduce a nonnegligible number of outliers in the selected samples because of the low detection rate from the low number of images. We employed the same classical DS MLE to estimate the deformation rate in both cases. Therefore, any improvement was solely due to the use of a more robust covariance matrix estimate.

The test area is mainly vegetation, except in the center where there is a road that usually appears as a DS in the X-band images. We had expected homogeneous deformation rates as the spans of the test area was roughly 100 m. However, many estimates appear in the left-hand figures as salt-and-pepper



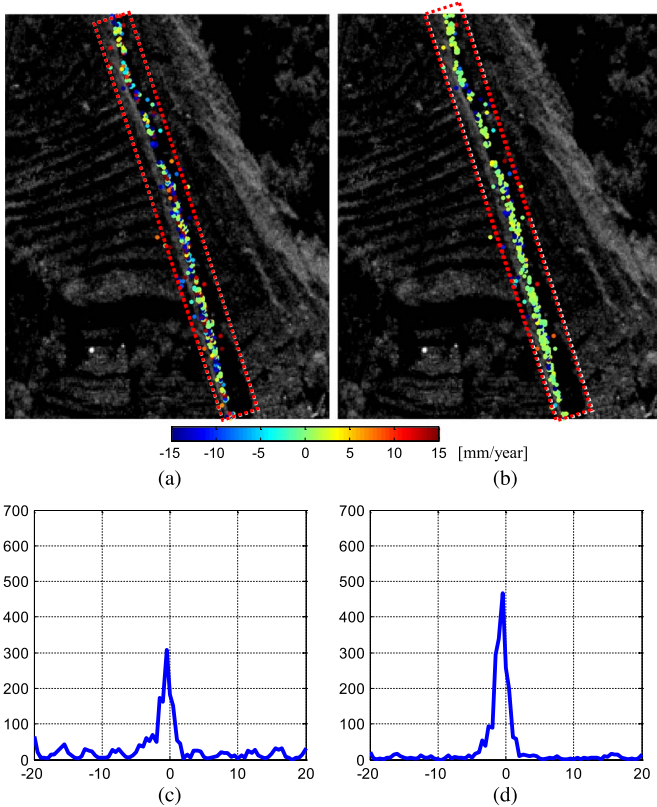


Fig. 4. Comparison of the linear deformation rate of the two test sites estimated using  $\hat{C}_{MLE}$  and  $\hat{C}_{RME}$ . Results (overlaid on the SAR intensity images) are specific to a local reference point. A homogeneous deformation rate is expected because of the small size of the area. (a) Results using the classical  $\hat{C}_{MLE}$ . (b) Results using the proposed  $\hat{C}_{RME}$ . The samples were identical for both covariance matrix estimations. They were adaptively selected (KS test) using *ten* images, which was expected to introduce a nonnegligible number of outliers in the selected samples because of the low detection rate from the low number of images. The same classical DS MLE was employed for the deformation rate estimations in both cases. (c) and (d) plot the corresponding histograms of the linear deformation rates enclosed in the two dashed red rectangles. In (c), many local peaks appear in the search range, most of which should not correspond to the deformation signals.

noise. We believe this is due to the low detection rate of the KS test and the nonstationarity of the samples. The improvement was clearly demonstrated.  $\hat{C}_{RME}$  significantly outperforms the widely used sample covariance matrix.

For further comparison, we plotted the histograms of the deformation rates enclosed in the two dashed red rectangles of Fig. 4(a) and (b). When using  $\hat{C}_{MLE}$ , many local peaks in the deformation rate appear in the search range. These peaks should not correspond to the deformation signal. With  $\hat{C}_{RME}$ , results are considerably more homogeneous, and thus more reasonable.

**2) Robustness Against Unmodeled Phase:** To evaluate the improvements by the robust loss function, we reintroduced the atmospheric phase of *ten* images randomly selected out of 33 back to the corrected phase to simulate realistic atmospheric phase error in 30% of the images. The results estimated by classical PS and DS phase parameter estimators are compared with those using their robust versions. The loss function of the robust estimators was chosen to be the Tukey's loss function with the  $c$  parameter set to 4.586. We also compare the difference of using

sample covariance matrix  $\hat{C}_{MLE}$  and rank covariance matrix  $\hat{C}_{RME}$  in the robust DS estimator.

Fig. 5(a) and (b) compare the linear deformation rates of the combined PSs and DSs in a different test area, which are estimated using the standard and robust estimators, respectively. The same sample covariance matrix  $\hat{C}_{MLE}$  was used in the estimation. The robust estimators outperform the classical estimators in regard of the variance of the estimates because, in this small area, we expect a homogeneous deformation rate. The improvement is solely due to the use of the robust loss function. In Fig. 5(c), we replaced the  $\hat{C}_{MLE}$  used in Fig. 5(b) with the proposed  $\hat{C}_{RME}$ . More improvement can be observed. It demonstrates that both phase error and poor sample selection can significantly degrade the performance of classical PS and DS MLEs. However, the results can be significantly improved by introducing a robust loss function and using a better covariance matrix estimate.

For reference, Fig. 5(d) shows the results using standard PS+DS estimators with the atmospheric phase of all image compensated. Under 30% of phase contamination, the robust estimators are able to achieve comparable result to the standard estimators with no phase contamination.

**3) Large Area Processing:** To assess overall performance, we applied our robust estimators over the entire image area of super volcano Campi Flegrei. Fig. 6(a) shows the combined results of the linear deformation rate using the proposed RIO framework, in which Tukey's loss function is used for both robust PS and DS estimator. We specifically chose the SCM for the robust covariance estimate because of the absence of significant fringes in the interferogram. With this combination, the proposed method retrieved 15 times more scatterers than that using only PS. This number should also be valid for standard PS+DS methods, e.g., SqueeSAR. However, RIO improves the overall robustness. This can be seen in the small area comparison [see Fig. 6(b)] of the two areas marked by red rectangles in Fig. 6(a), where RIO greatly reduced the salt-and-pepper noise probably caused by unmodeled phase and bad covariance estimation.

## VI. CONCLUSION AND OUTLOOK

The proposed RIO framework answers two open questions with respect to current multipass InSAR techniques: 1) how to downweight images with large phase error, such as the uncompensated atmospheric phase; and 2) how to correctly estimate the covariance matrix from nonstationary samples. We have provided corresponding solutions by introducing a robust loss function in the estimation of the phase history parameters and by using the *rank* of the complex multivariate together with robust weights (RME) in the covariance matrix estimation.

We demonstrate the benefits of the proposed framework by applying it to state-of-the-art estimators of PS and DS phase history parameters. Based on a simulation with 40% of its images being random-phase contaminated, the robust loss function increases the efficiency of the estimator by a factor of 7–35 compared with standard version estimators. The rank covariance  $M$ -estimator also tackles nonstationary samples to



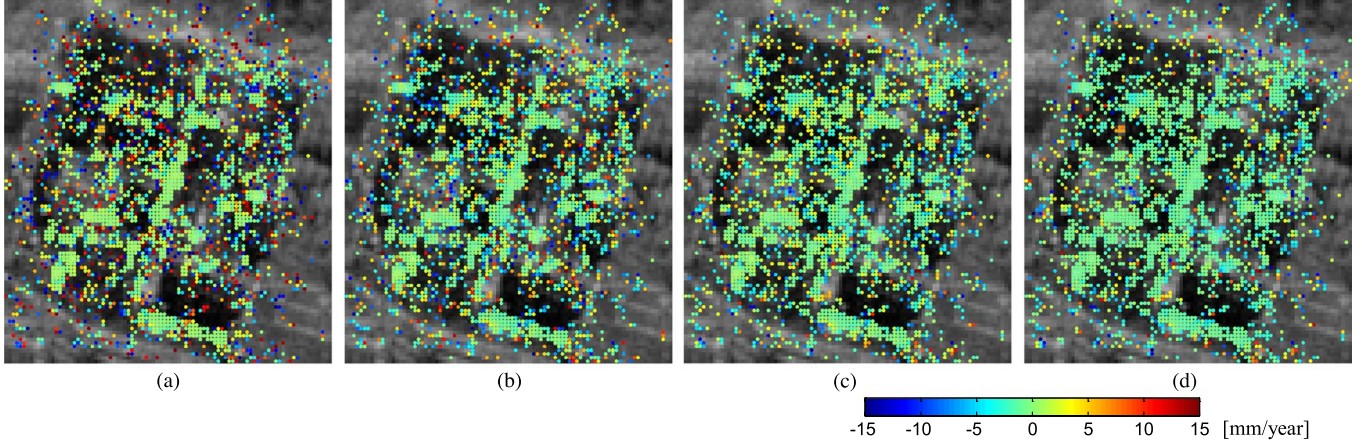


Fig. 5. Comparison of the linear deformation rate estimated using (a) standard PS and DS estimators with the sample covariance matrix  $\hat{\mathbf{C}}_{\text{MLE}}$  used in the DS estimator, (b) the proposed robust PS and DS phase parameter estimators using  $\hat{\mathbf{C}}_{\text{MLE}}$ , and (c) the proposed robust PS and DS phase parameter estimators using the rank of covariance matrix  $\hat{\mathbf{C}}_{\text{RME}}$ . The atmospheric phases of ten images out of 33 are not compensated, in order to simulate certain phase error. The loss functions of both the robust PS and DS estimators were chosen to be the Tukey's loss function with the  $c$  parameter set to 4.586. (d) shows the results using standard PS and DS estimators with all the atmospheric phase compensated. Under 30% of phase contamination, the proposed RIO method are able to achieve comparable result to the standard estimators with no phase contamination.

achieve results well beyond those of existing methods. For stationary samples, we propose alternative covariance estimators, i.e., an iterative  $M$ -estimator of covariance and a noniterative version for fast processing: sign covariance matrix.

The proposed framework can be easily extended to other multipass InSAR techniques without requiring excessive modification to the current programming. The computational effort will be approximately 3 to 4 times more than that required by standard processing, largely due to using the iterative method for solving the robust loss function, which usually converges within several iterations.

## APPENDIX

### A. Expectation for Rank Covariance of the SAR Acquisitions

Let us assume  $g_1$  and  $g_2$  are two stationary CCG random variables with covariance  $\gamma$ , and  $g_3$  and  $g_4$  are their direct neighborhood samples that have the same covariance  $\gamma$ , respectively. For simplicity, we suppose that all the random variables are normalized, i.e.,  $\sigma^2 = 1$ , so that  $\gamma \in [0, 1]$ , which is the coherence commonly used in SAR.

Then,  $g_1$  and  $g_2$  can be decomposed into three mutually uncorrelated stationary circular white Gaussian processes  $a$ ,  $b$ , and  $c$ , of equal variance, i.e.,

$$\begin{aligned} g_1 &= \sqrt{1-\gamma}a + \sqrt{\gamma}c \\ g_2 &= \sqrt{1-\gamma}b + \sqrt{\gamma}c. \end{aligned} \quad (31)$$

The same holds for  $g_3$  and  $g_4$ , i.e.,

$$\begin{aligned} g_3 &= \sqrt{1-\gamma}d + \sqrt{\gamma}f \\ g_4 &= \sqrt{1-\gamma}e + \sqrt{\gamma}f. \end{aligned} \quad (32)$$

The rank covariance/coherence of  $g_1$  and  $g_2$ , as defined in this paper (using one neighborhood sample), is then given as follows:

$$E\{(g_1 g_3^*)(g_2 g_4^*)\} = E\{(\gamma c f^*)(\gamma f c^*)\} = \gamma^2. \quad (33)$$

### B. Weighting Function of Complex Multivariate $t$ -Distribution

In general, the weighting function can be obtained by following (7). However, this method requires the transformation of the complex  $t$ -distribution into the real domain so that the derivation can be performed. Therefore, we formed a new real multivariate  $\tilde{\mathbf{g}}$  using the real and imaginary parts of the complex multivariate, i.e.,

$$\begin{aligned} \tilde{\mathbf{g}} &= \begin{bmatrix} \text{Re}(\mathbf{g}) \\ \text{Im}(\mathbf{g}) \end{bmatrix} \in \mathbb{R}^{2N} \\ \mathbf{C}_{\tilde{\mathbf{g}}\tilde{\mathbf{g}}} &= \frac{1}{2} \begin{bmatrix} \text{Re}(\mathbf{C}_{\mathbf{g}\mathbf{g}}) & -\text{Im}(\mathbf{C}_{\mathbf{g}\mathbf{g}}) \\ \text{Im}(\mathbf{C}_{\mathbf{g}\mathbf{g}}) & \text{Re}(\mathbf{C}_{\mathbf{g}\mathbf{g}}) \end{bmatrix} \in \mathbb{R}^{2N \times 2N} \end{aligned} \quad (34)$$

$\tilde{\mathbf{g}}$  follows a  $2N$ -variate real  $t$ -distribution with the covariance matrix being  $\mathbf{C}_{\tilde{\mathbf{g}}\tilde{\mathbf{g}}}$ , i.e.,

$$f_{\tilde{\mathbf{g}}}(\tilde{\mathbf{g}}) = \frac{\Gamma(\frac{\nu+2N}{2})}{(\pi\nu)^N \Gamma(\frac{\nu}{2}) \det(\mathbf{C}_{\tilde{\mathbf{g}}\tilde{\mathbf{g}}})^{\frac{1}{2}}} \left(1 + \frac{1}{\nu} \tilde{\mathbf{g}}^T \mathbf{C}_{\tilde{\mathbf{g}}\tilde{\mathbf{g}}}^{-1} \tilde{\mathbf{g}}\right)^{-\frac{\nu+2N}{2}} \quad (35)$$

which, according to [33], is equivalent to the complex  $t$ -distribution.

Let the squared residual  $\tilde{x}^2 = \tilde{\mathbf{g}}^T \mathbf{C}_{\tilde{\mathbf{g}}\tilde{\mathbf{g}}}^{-1} \tilde{\mathbf{g}}$ . As  $\tilde{x}$  is whitened, it follows a univariate  $t$ -distribution, i.e.,

$$f_{\tilde{x}}(\tilde{x}) = C(1 + \nu^{-1}\tilde{x}^2)^{-(\nu+2N)/2} \quad (36)$$

where  $C$  is the lengthy constant in the beginning of (34).  $C$  will disappear after considering the derivative. Now, the weighting function can be derived following (7), which is  $(2N + \nu)/(\nu + \tilde{x}^2)$ . Transforming it back to the complex domain, we introduce a factor of 2, as  $\tilde{\mathbf{g}}^T \mathbf{C}_{\tilde{\mathbf{g}}\tilde{\mathbf{g}}}^{-1} \tilde{\mathbf{g}} = 2\mathbf{g}^H \mathbf{C}_{\mathbf{g}\mathbf{g}}^{-1} \mathbf{g} \Rightarrow \tilde{x}^2 = 2x^2$ . Hence, the corresponding weighting function for complex  $t$ -distribution is  $(2N + \nu)/(\nu + 2x^2)$ .



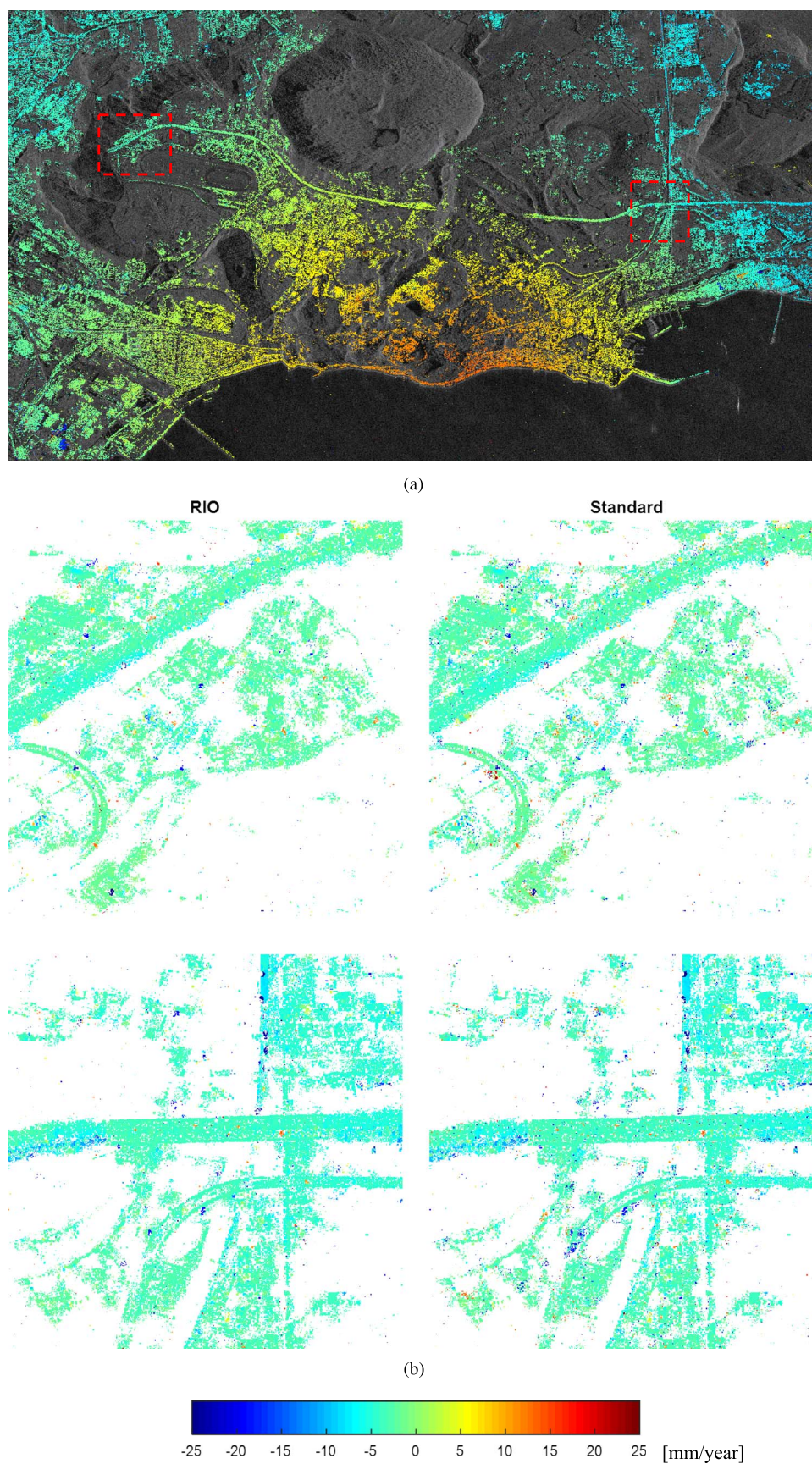


Fig. 6. (a) Linear deformation rate of the super volcano Campi Flegrei, as retrieved using the proposed RIO framework that combines both PS and DS in a stack of 33 TS-X high-resolution spotlight images. Tukey's biweight function is set to be the robust loss function, and the sign covariance matrix is employed as the robust covariance matrix estimate. It retrieves 15 times more scatterers than those using only PS only. This number should also be valid for standard PS+DS methods, e.g., SqueeSAR. However, RIO greatly reduced the salt-and-pepper noise probably caused by unmodeled phase and bad covariance estimation. This can be seen in the small area comparison in (b), which plots the results for the two areas marked by red rectangles in (a).

## ACKNOWLEDGMENT

The authors would like to thank Prof. R. Bamler for valuable discussion.

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**Yuanyuan Wang** (S'10) received the B.Eng. (Hons) degree in electrical engineering from The Hong Kong Polytechnic University, Hong Kong, in 2008 and the M.Sc. degree in Earth oriented space science and technology from Technische Universität München (TUM), Munich, Germany, in 2010. Since 2011, he has been pursuing the Doctoral degree with the International Graduate School of Science and Engineering, TUM, and currently with the project "Joy of Long Baseline" of German Aerospace Center.

In June and July 2014, he was a Guest Scientist with the Institute of Visual Computing, ETH Zürich, Zürich, Switzerland. He is currently with Signal Processing in Earth Observation (SiPEO), TUM. His research interests include optimal parameters estimation in interferometric SAR techniques, multisensor fusion algorithms of SAR/optical data, and the applications of these techniques in urban and volcanic areas.



**Xiao Xiang Zhu** (S'10–M'12–SM'14) received the Bachelor's degree in space engineering from the National University of Defense Technology, Changsha, China, in 2006 and the M.Sc. degree, Dr.-Ing. degree, and Habilitation in signal processing from Technische Universität München (TUM), Munich, Germany, in 2008, 2011, and 2013, respectively.

She was a Guest Scientist or a Visiting Professor with the Italian National Research Council (CNR-IREA), Naples, Italy, in 2009; with Fudan University, Shanghai, China, in 2014; and with the University of Tokyo, Tokyo, Japan in 2015.

Since 2011, she has been a Scientist with the Remote Sensing Technology Institute, German Aerospace Center (DLR), Oberpfaffenhofen, where she is currently the Head of the Signal Analysis Team. Since 2013, she has also been a Helmholtz Young Investigator Group Leader and appointed as a TUM Junior Fellow. She is currently a Professor of signal processing in earth observation with TUM. Her main research interests include advanced interferometric SAR techniques such as high-dimensional tomographic SAR imaging and SqueeSAR; computer vision in remote sensing such as object reconstruction and multidimensional data visualization; and modern signal processing, such as innovative algorithms (e.g., compressive sensing and sparse reconstruction), with applications in the field of remote sensing such as multi/hyperspectral image analysis.

Dr. Zhu serves as an Associate Editor for IEEE TRANSACTIONS ON GEOSCIENCE AND REMOTE SENSING.





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